

# Mapping local patterns of childhood overweight and wasting in low- and middle-income countries between 2000 and 2017

**LBD Double Burden of Malnutrition Collaborators**

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# Mapping local patterns of childhood overweight and wasting in low- and middle-income countries between 2000 and 2017

LBD Double Burden of Malnutrition Collaborators\*

**A double burden of malnutrition occurs when individuals, household members or communities experience both undernutrition and overweight. Here, we show geospatial estimates of overweight and wasting prevalence among children under 5 years of age in 105 low- and middle-income countries (LMICs) from 2000 to 2017 and aggregate these to policy-relevant administrative units. Wasting decreased overall across LMICs between 2000 and 2017, from 8.4% (62.3 (55.1–70.8) million) to 6.4% (58.3 (47.6–70.7) million), but is predicted to remain above the World Health Organization's Global Nutrition Target of <5% in over half of LMICs by 2025. Prevalence of overweight increased from 5.2% (30 (22.8–38.5) million) in 2000 to 6.0% (55.5 (44.8–67.9) million) children aged under 5 years in 2017. Areas most affected by double burden of malnutrition were located in Indonesia, Thailand, southeastern China, Botswana, Cameroon and central Nigeria. Our estimates provide a new perspective to researchers, policy makers and public health agencies in their efforts to address this global childhood syndemic.**

The profound impacts of childhood malnutrition, including both undernutrition and overweight, affect the economic, social and medical well-being of individuals, families, communities and nations<sup>1,2</sup>. Undernutrition has been the most common form of malnutrition in LMICs<sup>3</sup>, but as populations experience economic growth, urbanization and demographic change, overweight is an emerging problem, leading to a double burden of malnutrition (DBM). DBM may be manifested at the individual level as stunting in childhood followed by overweight in adulthood<sup>4</sup>. At the household level, research has focused on maternal and child indicators of malnutrition, whereas at the population level, prevalence of both undernutrition with overweight has been reported<sup>5</sup>. In children, DBM can be defined using different combinations of the various indicators of undernutrition (wasting and/or stunting) and overweight, obesity and diet-related noncommunicable diseases (NCDs)<sup>6</sup>. While the most studied type of double burden is that of stunting and obesity, it is mostly applicable at the individual level among overweight adults who were previously stunted from chronic undernutrition during childhood. Wasting is associated with high rate of child mortality, whereas stunting has significant negative impact across the life course and is highly predictive of economic outcomes<sup>7</sup>. Public health nutrition programs designed to address undernutrition may exacerbate overweight<sup>8</sup>, thus a comprehensive understanding of DBM at the population level is crucial for the design of effective interventions.

Our aim was to determine the prevalence of overweight among children under 5 years old in LMICs ( $N = 105$ ) for policy-relevant administrative units (district, state, and national level) and determine DBM by combining these estimates with those of wasting prevalence. As there is no broad consensus on the preferred international child growth standards for assessing overweight and obesity among children under 5 (refs. <sup>9,10</sup>), we used weight-for-height above established cutoff points defined by the World Health Organization (WHO). This was to analyze overweight estimates in relation to the Global Nutrition Targets (GNTs), which were developed based on WHO standards. Prevalence of early childhood overweight

(including obesity) is defined as the proportion of children under 5 with a weight-for-height  $z$  score (WHZ) more than two standard deviations (s.d.) above the WHO sex- and age-specific median growth reference standards<sup>10</sup>. This is different from the definition for children between the ages of 5–18 years, which is above one s.d. for overweight and above two s.d. for obese. We selected wasting as the comparative indicator against overweight, as both share recommended population prevalence ranges, which can be used to create bivariate categories for DBM. Child wasting prevalence is defined as the proportion of children under 5 with a WHZ more than two s.d. below the median WHO growth standards<sup>10</sup>. Using WHZs allowed modeling of the three categories in the same distribution and thus enabled us to reliably determine the relative proportions for each category using an ordinal approach. Based on WHO and United Nations Children's Fund (UNICEF)-defined thresholds, a moderate level of separate or dual conditions is defined as >5–10%, a high level as >10–15% and a very high level as >15% estimated prevalence<sup>11</sup>. Finally, we have defined DBM in this study as the simultaneous occurrence of >5% estimated prevalence for both wasting and overweight within the same locations in the same year.

Reversing the rise in childhood overweight is indicated in the United Nations (UN) Sustainable Development Goal 2.2 (ref. <sup>12</sup>) and WHO's GNTs to improve maternal, infant and young child nutrition<sup>13</sup>. WHO has also set an international target to reduce wasting to <5% by 2025 (ref. <sup>14</sup>). Quantifying changes in childhood overweight and wasting prevalence can be used to measure progress toward these targets, while identifying locales with simultaneous overweight and wasting will better inform intervention planning. In addition, mapping changes in DBM prevalence will provide a deeper understanding of the impact of past intervention strategies, including insight into overweight in children under 5.

## Global and local variation in malnutrition trends

Globally in 2017, an estimated 38.3 million (5.6%) children under 5 were overweight and 50.5 million (7.5%) were wasted<sup>15</sup>. The majority (91%) of children under 5 affected by wasting and nearly half

\*A list of authors and their affiliations appears online. ✉e-mail: [sihay@uw.edu](mailto:sihay@uw.edu)

(48%) of overweight children lived in LMICs, with Africa and Asia accounting for the largest shares of the global burden (25% and 46% of overweight and 27% and 69% of wasted children, respectively)<sup>16</sup>. Direct comparisons of population-level trends of childhood overweight and wasting generally provide regional- or country-level estimates<sup>5,16–20</sup>, potentially masking important subnational differences. Previously, we mapped 2000–2017 prevalence and trends in wasting, stunting and underweight among children under 5 across LMICs<sup>21</sup> using Bayesian model-based geostatistical techniques<sup>22</sup>. Building from this approach and using data from 420 household surveys representing more than 3 million children, we mapped the relative burdens of overweight and wasting among children under 5 in 105 LMICs from 2000 to 2017. Mapping with a continuous model allows us to incorporate geolocated data and covariates and produce gridded cell-level estimates that can be aggregated to intervention- or policy-relevant geographical areas as boundaries change over time. We present estimates at this local grid cell-level and aggregate to first administrative (such as states and provinces), second administrative (such as districts and departments) and national levels. On the basis of 2000 to 2017 weighted annualized rates of change (AROC), which apply more weight to recent data, we predict prevalence of overweight and wasting and estimate their double burden in 2025. The full array of outputs are available at the Global Health Data Exchange (<http://ghdx.healthdata.org/record/ihme-data/lmic-double-burden-of-malnutrition-geospatial-estimates-2000-2017>) and can be further explored with our customized visualization tools (<https://vizhub.healthdata.org/lbd/dbm>).

### Prevalence and trends in early childhood overweight

Across LMICs, the prevalence of early childhood overweight increased from 5.2% (95% uncertainty interval, 4.5–5.4%) to 6.0% (4.8–6.1%) in the modeled study period. Between 2000 and 2017, there were noticeable differences in estimated levels by area (Fig. 1a,b). Although levels varied broadly across LMICs, every modeling region had areas with high estimated prevalence in 2017 (Fig. 1b and Extended Data Fig. 1). These included large contiguous areas across most Central American, Caribbean and South American countries and areas with  $\geq 15\%$  estimated prevalence in central Cuba, southern Panama, western Paraguay, scattered throughout several eastern Brazilian states (for example, in Rio Grande do Sul, Minas Gerais, Santa Catarina, Paraná and São Paulo) and Peru's coastal cities of Tacna, Ilo, Islay, Callao, Trujillo and Lima. In Africa, most countries bordering the Sahel had low overweight prevalence (0–5%); areas with  $>15\%$  estimated prevalence were concentrated in North Africa throughout Morocco, Algeria, Tunisia, Egypt and select areas of Libya, as well as along South Africa's southern coast and in pockets in Botswana and Zambia. Large areas in eastern and northern China and throughout Mongolia had an estimated overweight prevalence  $>15\%$ . Countries in the Oceania region had moderate to high levels, with estimates over 15%, such as in Indonesia's Jakarta Pusat and Jakarta Barat regencies (in Jakarta Raya; 17.7% (15.3–18.4%)). The North Africa, Central Asia and Southeast Asia regions showed vast differences across nations; for example, Afghanistan, Sudan and Laos had  $<5\%$  estimated national prevalence, whereas Egypt, Uzbekistan, Morocco, Kyrgyzstan and Thailand had  $\geq 15\%$ . South Asia's estimated levels ranged from  $<5\%$  in Bangladesh to  $\geq 10\%$  Bhutan. Estimated prevalence in Karbala city in Karbala, Iraq, increased from 13.6% (12.4–14.1%) in 2000 to 29.3% (22.9–29.1%) in 2017. Thailand's southern areas experienced large increases in estimated prevalence levels; Sathorn district, Bangkok Metropolis, had 24.1% (20.1–24.8%) overweight in 2000 and 33.9% (27.5–35.5%) in 2017. Areas with the greatest decrease included Churcampa district, Huancavelica, Peru, decreasing from 17.5% (17.4–17.6%) in 2000 to 10.3% (10.2–10.4%) in 2017. Similarly, overweight in Al Gash district, Kassala, Sudan, declined from 14.1% (13.6–14.5%) to 6.1% (5.2–6.2%).

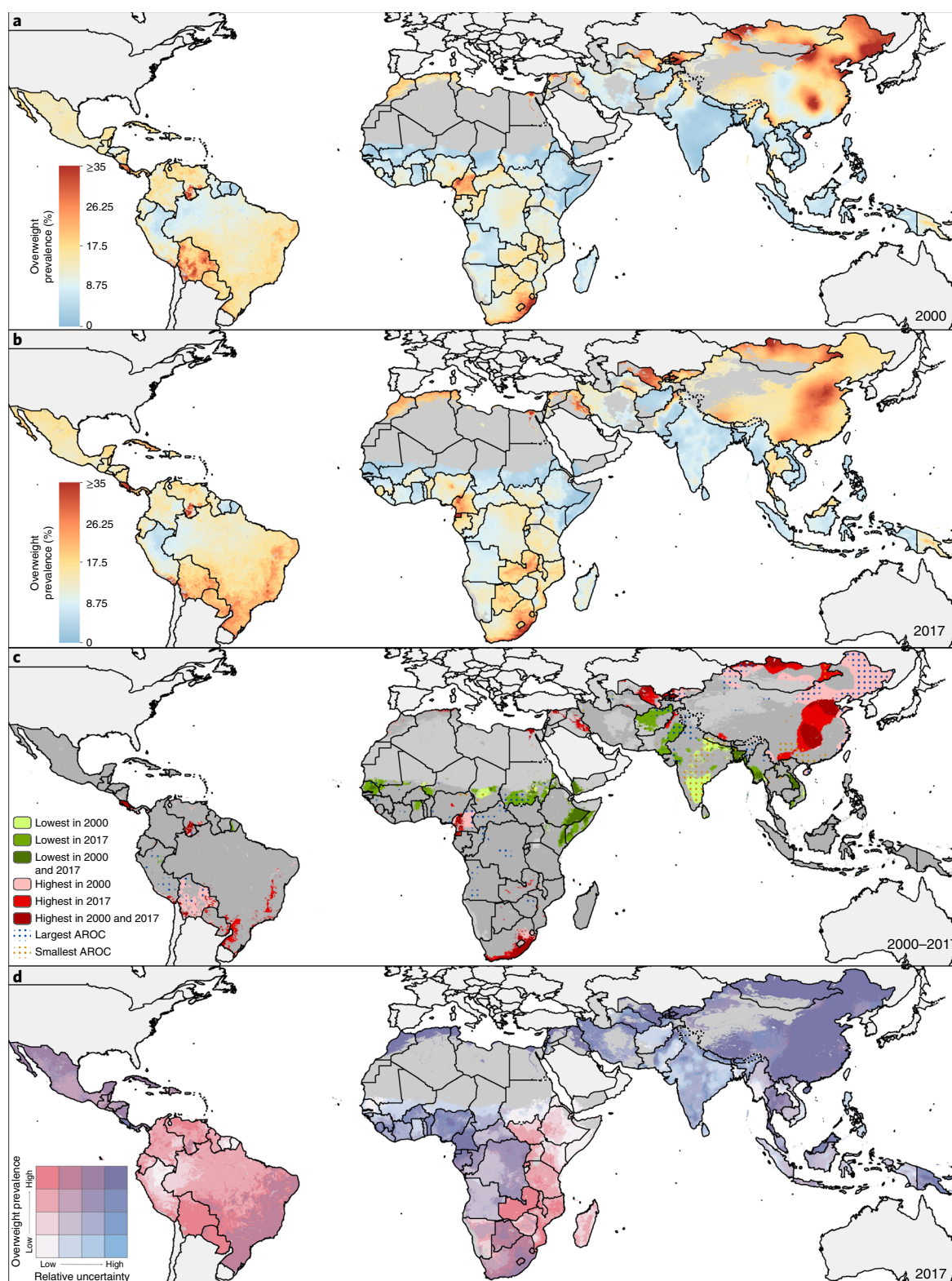
Within-country differences in estimated overweight levels were found in 37 (35.2%) LMICs, including South Africa, Peru and Indonesia, which had twofold differences in estimated prevalence across second administrative units in 2017. South Africa had high estimated national levels (24.9% (23.9–25.2%)); however, the province of Northern Cape had moderate levels (14.6% (13.6–14.9%)), whereas the southeastern province of Eastern Cape had very high levels (32.7% (30.8–33.9%)). Disparities were further pronounced at the district level. Siyanda (Northern Cape) had 12.5% (11.6–12.9%) prevalence, whereas Ugu (KwaZulu-Natal) had 36.7% (34.0–38.2%). Nearly every modeling region had areas with overweight prevalence that ranked among the highest decile in 2000, 2017 or both years (Fig. 1c).

Overall, the number of overweight children under 5 in LMICs also showed a significant increase from 30.0 million (22.8–38.5) to 55.5 million (44.8–67.9) in the study period (Fig. 2a,b). By 2017, 26.2 million (24.1–27.2 million; 36.0%) of those affected lived in eastern Asia, northern Africa or South America. An estimated 8.6% (8.5–9.9%) of first administrative units had fewer than 1,000 overweight children under 5, 47.5% (47.2–49.5%) had 1,000 to  $<10,000$ , 43.8% (40.6–44.3%) had 10,000 to  $<100,000$  and just 3.8% (3.7–3.9%) had 100,000 or more. Some areas, such as northern and central parts of Bolivia, experienced large annualized declines such that their ranking among the highest estimated prevalence decile in 2000 no longer applied in 2017. In contrast, a large area in India, south of the Tropic of Cancer, experienced large annualized increases in overweight; its ranking among the lowest prevalence decile in 2000 was not maintained in 2017. All modeled regions had areas that experienced average annualized increases of  $\geq 1\%$  in overweight prevalence (Fig. 2c). Unless current trajectories change, prevalence of overweight will continue to increase to 2025 (Fig. 2d).

### Prevalence and trends in child wasting

The estimated prevalence of early childhood wasting decreased overall across LMICs between 2000–2017, from 8.4% (7.9–9.9%) to 6.4% (4.9–7.9%). The most notable relative reductions were seen across North Africa and in select countries in sub-Saharan African (SSA) regions, Central and Andean America and Southeast Asia regions. In Burkina Faso's Ganzourgou district, estimated levels declined from 20.2% (19.1–21.3%) in 2000 to 11.6% (10.9–12.1%) in 2017, in Yemen's Ash Shaikh Outhman district from 25.1% (22.2–26.3%) to 21.3% (18.9–22.2%) and in Sudan's Al Mahagil district from 31.9% (31.4–32.6%) to 12.2% (10.5–12.9%). Increases in estimated prevalence also occurred, such as in Pakistan's Makran district (Baluchistan), from 7.4% (6.7–7.6%) to 11.4% (10.4–11.8%).

In 2017, there were several instances of contrasting geographic patterns of child wasting compared to those of overweight. Many Central American, Caribbean and South American countries (46%; 11 of 24) affected by overweight ( $>15\%$  prevalence) met the WHO GNTs for  $<5\%$  prevalence of wasting across all districts based on estimated prevalence (Fig. 3a,b and Extended Data Fig. 2). Estimated wasting prevalence was  $\geq 15\%$  in 31.9% (850 of 2,661) and  $\geq 20\%$  in 12.9% (342) of second administrative units across Central and South Asian countries, contributing to high prevalence at the national level in India (15.7% (15.4–15.9%)), Pakistan (12.2% (11.8–12.4%)) and Sri Lanka (11.2% (10.5–11.5%)); Afghanistan and Bangladesh maintained high levels (estimated prevalence  $\geq 10\%$ ) across many areas. Local-level estimates delineate very high wasting prevalence ( $\geq 15\%$ ) along the African Sahel from Mauritania to Sudan, in the northeastern Horn of Africa and neighboring countries of Eritrea, Ethiopia, Somalia, Kenya, South Sudan and Yemen, in select areas in Algeria and Egypt, and across Madagascar. In the Middle East, Syria exceeded 15% estimated prevalence throughout most areas and Iraq's southeastern districts exceeded 10%. Estimated levels of wasting were relatively uniform and low across East Asia, with the exception of a few focal areas exceeding 10% or 20% in central

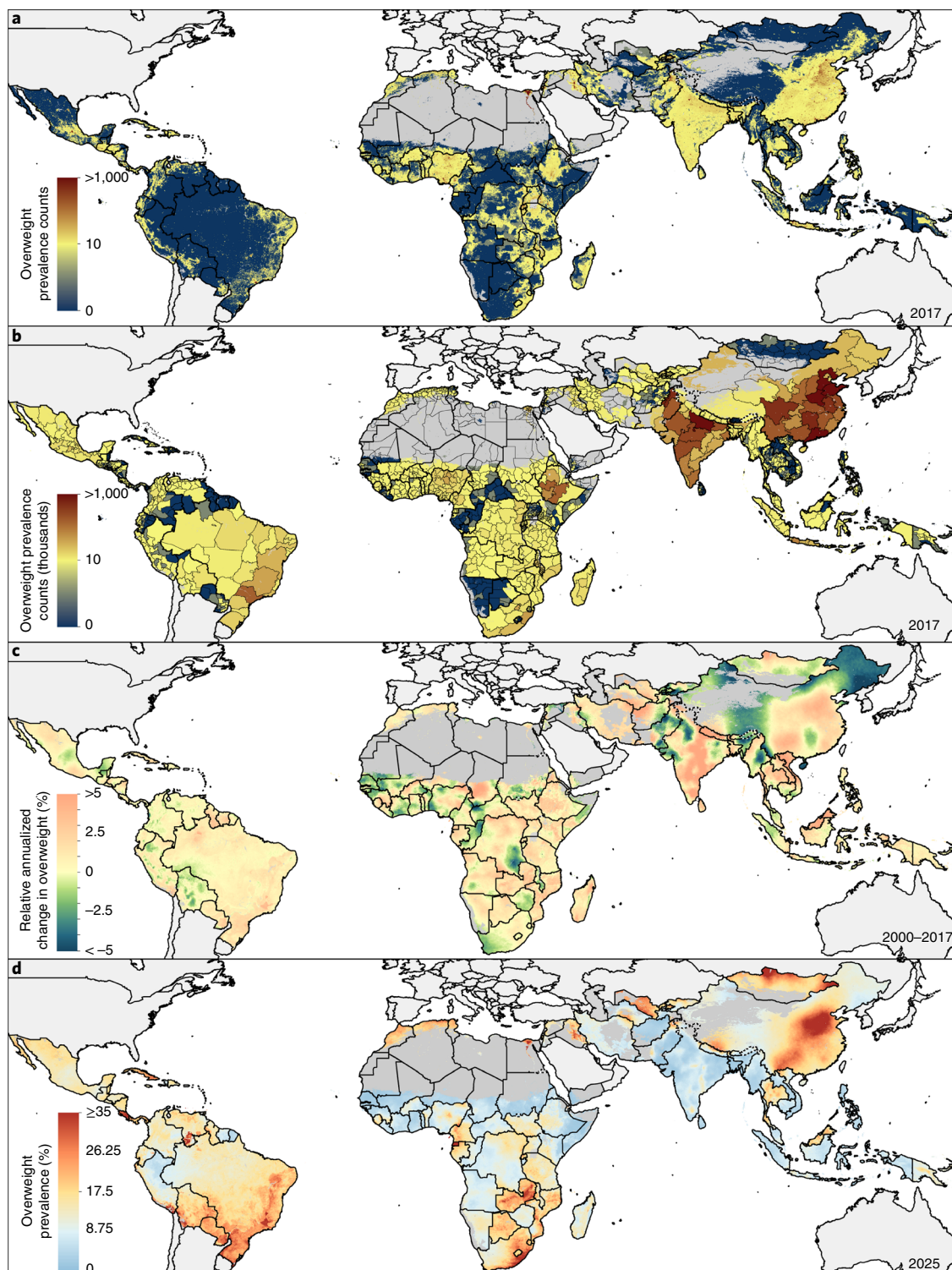


**Fig. 1 | Prevalence of overweight children under 5 in LMICs (2000–2017).** **a,b**, Prevalence of overweight among children under 5 at 5 × 5-km resolution in 2000 (**a**) and 2017 (**b**). **c**, Overlapping population-weighted lowest and highest 10% of grid cells and AROC in overweight from 2000 to 2017. **d**, Overlapping population-weighted quartiles of overweight and relative 95% uncertainty in 2017. Maps reflect administrative boundaries, land cover, lakes and population; gray colored areas have grid cells classified as ‘barren or sparsely vegetated’ and had fewer than ten people per 1 × 1-km grid cell in 2017 or were not included in this analysis<sup>39–45</sup>. Maps were generated using ArcGIS Desktop 10.6.

pockets of east China. Most areas in Southeast Asia and Oceania experienced moderate-to-high estimated wasting levels (~10%), whereas some areas in Indonesia’s southern-most islands in Nusa

Tenggara (Timor state) exceeded 15% prevalence. Meanwhile, some areas in Myanmar, Thailand, northern Laos and Vietnam had very low levels, approaching the WHO GNTs.

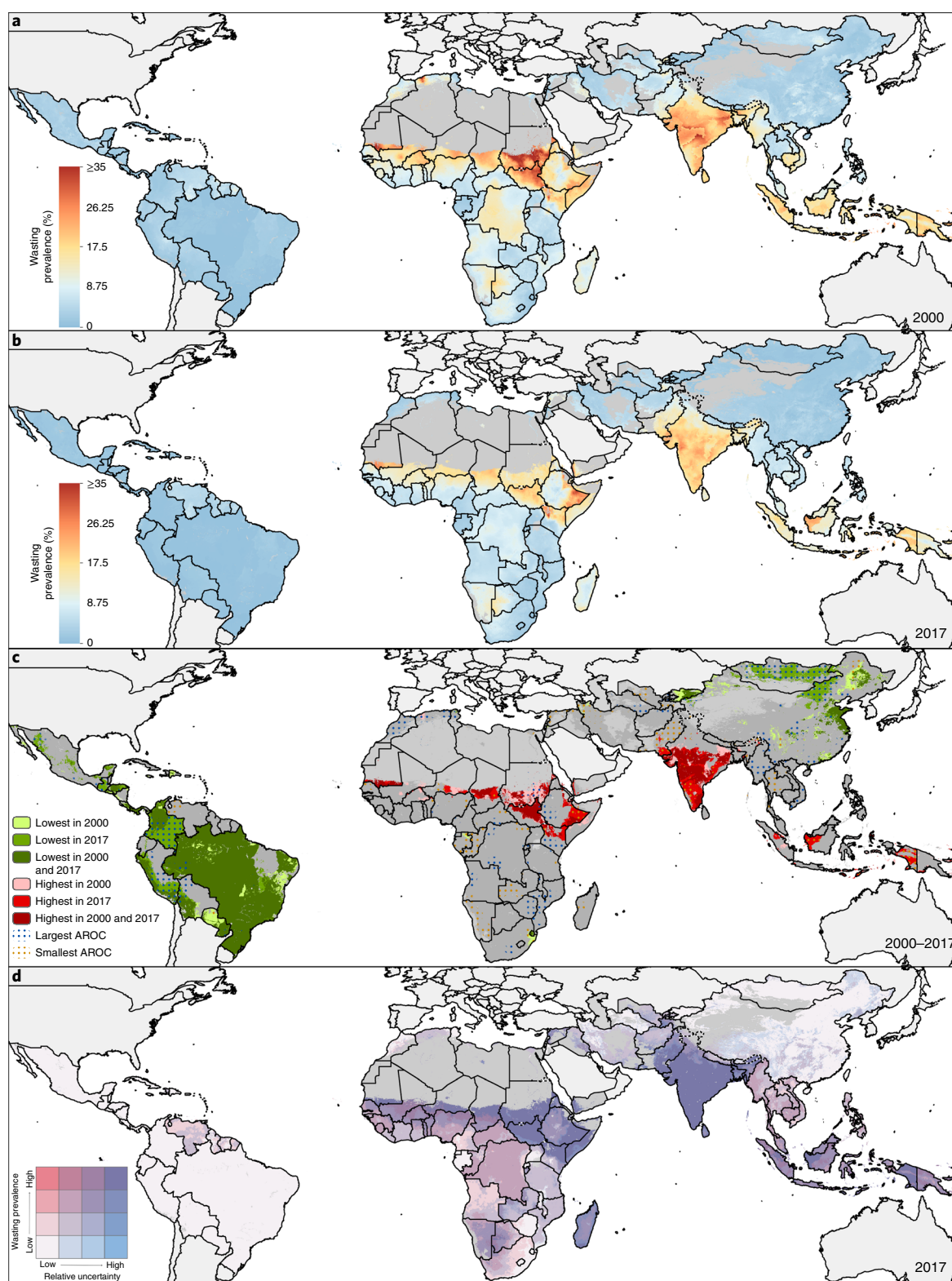




**Fig. 2 | Number of overweight children under 5 in LMICs (2000–2017) and progress toward 2025. a,b,** Number of children under 5 affected by overweight at a 5×5-km resolution (**a**) and by first administrative units (**b**). **c,** Annualized decrease (AD) in overweight prevalence from 2000 to 2017. **d,** Grid cell-level predicted overweight prevalence in 2025 based on AD achieved from 2000 to 2017 and projected from 2017. Maps reflect administrative boundaries, land cover, lakes and population; gray colored areas have grid cells classified as 'barren or sparsely vegetated' and had fewer than ten people per 1×1-km grid cell in 2017 or were not included in this analysis<sup>39–45</sup>. Maps were generated using ArcGIS Desktop 10.6.

Between 2000 and 2017, the number of children under 5 affected by wasting decreased from 62.3 (55.1–70.8) million to 58.3 (47.6–70.7) million, 28.4% (28.2–28.5) of whom were in Africa and 65.4% (63.6–67.3) in South Asia in 2017 (Fig. 3c,d). Despite maintaining high estimated prevalence in many areas, all regions

in Africa had areas that experienced among the highest rates of annualized declines in 2000–2017; only a few areas in Chad, Sudan, South Sudan, Ethiopia and Kenya were among the highest decile of estimated prevalence levels in both 2000 and 2017 (Fig. 4a,b). Progress differed across and within African countries, with some

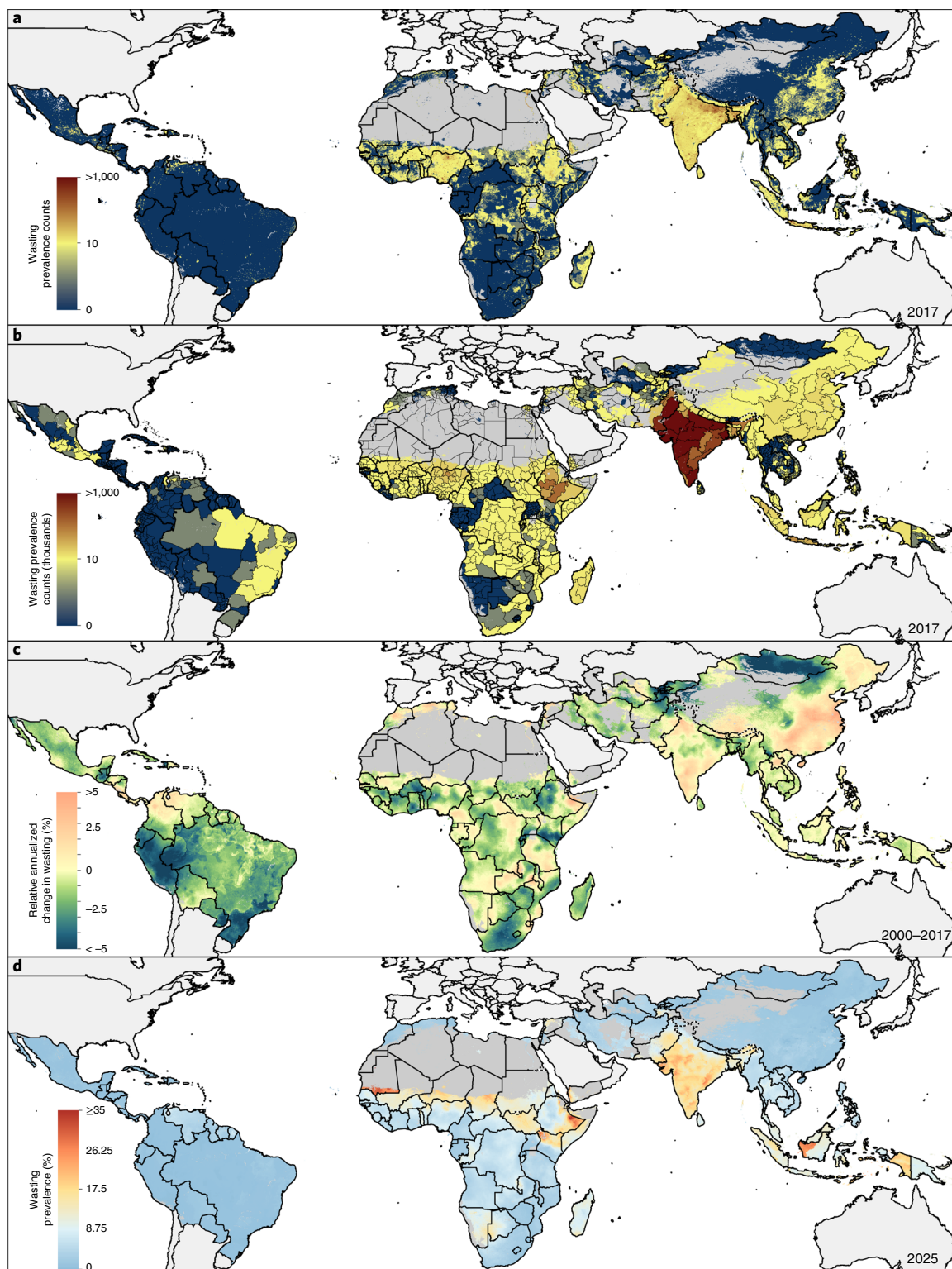


**Fig. 3 | Prevalence of wasted children under 5 in LMICs (2000–2017).** **a–c**, Prevalence of moderate and severe wasting among children under 5 at a 5 × 5-km resolution in 2000 (**a**) and 2017 (**b**). **c**, Overlapping population-weighted lowest and highest 10% of grid cells and AROC in wasting from 2000 to 2017. **d**, Overlapping population-weighted quartiles of wasting and relative 95% uncertainty in 2017. Maps reflect administrative boundaries, land cover, lakes and population; gray colored areas have grid cells classified as ‘barren or sparsely vegetated’ and had fewer than ten people per 1 × 1-km grid cell in 2017 or were not included in this analysis<sup>39–45</sup>. Maps were generated using ArcGIS Desktop 10.6.

nations, such as Nigeria, Ethiopia and Namibia, experiencing both annualized decreases and increases in wasting within their borders (Fig. 4c). Overall, South America and South SSA demonstrated

the largest annualized declines ( $\geq 5\%$ ) across most of their areas and regions of Latin America and the Caribbean, the Middle East, South Asia, Southeast Asia and Oceania experienced mostly





**Fig. 4 | Number of wasted children under 5 in LMICs (2000–2017) and progress toward 2025. a,b,** Number of children under 5 affected by wasting at the 5×5-km resolution (**a**) and by first administrative units (**b**). **c,** AD in wasting prevalence from 2000 to 2017. **d,** Grid cell-level predicted wasting prevalence in 2025 based on AD achieved from 2000 to 2017 and projected from 2017. Maps reflect administrative boundaries, land cover, lakes and population; gray colored areas have grid cells classified as 'barren or sparsely vegetated' and had fewer than ten people per 1×1-km grid cell in 2017 or were not included in this analysis<sup>39–45</sup>. Maps were generated using ArcGIS Desktop 10.6.

annualized increases. Large areas of India and parts of central Pakistan experienced some of the highest prevalence levels throughout the study period, as well as annualized increases. Nearly all South Asian countries had large contiguous areas of stagnation or

annualized increases in wasting; given recent rates of progress, few will meet the WHO GNTs in all their locations by 2025 (Fig. 4d). By 2025, 68 (64.8%) of LMICs are predicted to fail to meet the <5% target nationally, all of which are in Africa, Asia and the Middle East.

Based on subnational estimates, 88 (83.8%) and 94 (89.5%) will fail to meet the wasting WHO GNTs in all first and second administrative units, respectively.

### Double burden of wasting and overweight

Nearly every modeling region had subnational areas with at least moderate co-occurrence of wasting and overweight ( $\geq 5\%$  estimated prevalence of both conditions) in 2017 (Fig. 5 and Extended Data Fig. 3). Exceptions were Central and South America, where Guyana was the only example of moderate DBM (5%–10% of both conditions). In Africa, much of the Democratic Republic of the Congo, Cameroon, Republic of Congo, Zambia and southern Botswana demonstrated high DBM ( $\geq 10\%$  of both overweight and wasting). Areas in central Morocco reached some of the highest levels of DBM ( $\geq 15\%$  overweight, 10–15% wasting), whereas much of the rest of North Africa had high estimated overweight (10–15%) and moderate estimated wasting (5–10%). Locations scattered throughout Iraq, India and in Southeast Asia mostly experienced moderate wasting (such as Myanmar at 5–10%) or moderate DBM (such as Indonesia at 5–10%), reaching moderate-to-high DBM levels in select areas (such as central Papua New Guinea and Cambodia at 5–10% overweight, 10–15% wasting; Thailand, 10–15% overweight, 5–10% wasting). Relatively rare in East Asia, DBM was at moderate levels at most (5–10% both conditions), such as in provinces in southeastern China. At the national level, 25.7% (27 of 105) LMICs were moderately affected and 5.7% (6 of 105) were highly affected by both overweight and wasting ( $\geq 5\%$  and  $\geq 10\%$  prevalence of both conditions, respectively). Subnationally, however, 70.5% (74 of 105) of LMICs had moderately affected districts, 11.4% (12 of 105) had highly affected districts and 2.9% (3 of 105) had districts with very high DBM ( $\geq 5\%$ ,  $\geq 10\%$  and  $\geq 15\%$  prevalence of both conditions, respectively).

Although childhood nutritional status generally improved over 2000–2017, subnational variation in childhood overweight, wasting and DBM was apparent. Declines in wasting and overweight prevalence in South Africa's western areas led to a decrease in DBM prevalence, from high levels in Siyanda district in 2005 (10–15% estimated wasting and overweight) to moderate levels in 2017 (5–10% both conditions); overweight remains very high, however, on the southern coast ( $\geq 15\%$ ). On the basis of annualized trends, 25.7% (27 of 105) of LMICs are predicted to have districts with at least moderate DBM by 2025 and 34.3% (36 of 105) are predicted to have high DBM districts (Fig. 5). Between 2000 and 2017, 8.6% (9 of 105) of LMICs had first administrative units that experienced transition from high estimated prevalence of wasting ( $\geq 10\%$ ) to normal weight ( $< 5\%$  both wasting and overweight). Nearly one-third, 32.3% (34 of 105) of LMICs had first administrative units that transitioned from normal weight to high overweight and 7.6% (8 of 105) transitioned from high wasting to high DBM.

### Discussion

This study provides overweight estimates and combines them with wasting estimates to highlight DBM across LMICs at a fine geospatial scale. This enables efficient targeting of local-level interventions to improve nutrition outcomes in vulnerable populations. The figures presented here, as well as our online visualization tools, allow for comparing overweight and wasting levels and trends across and within countries for each year from 2000 to 2017, leveraging the spatially resolved underlying data and covariates to produce detailed spatial estimates across all modeled regions. Our estimates show the global trend in early childhood wasting is declining, but areas with high prevalence and little progress, such as in the Sahel and South Asia, remain. Meanwhile, childhood overweight prevalence has increased, especially in tropical South America and regions in the Middle East, Central Asia and Africa.

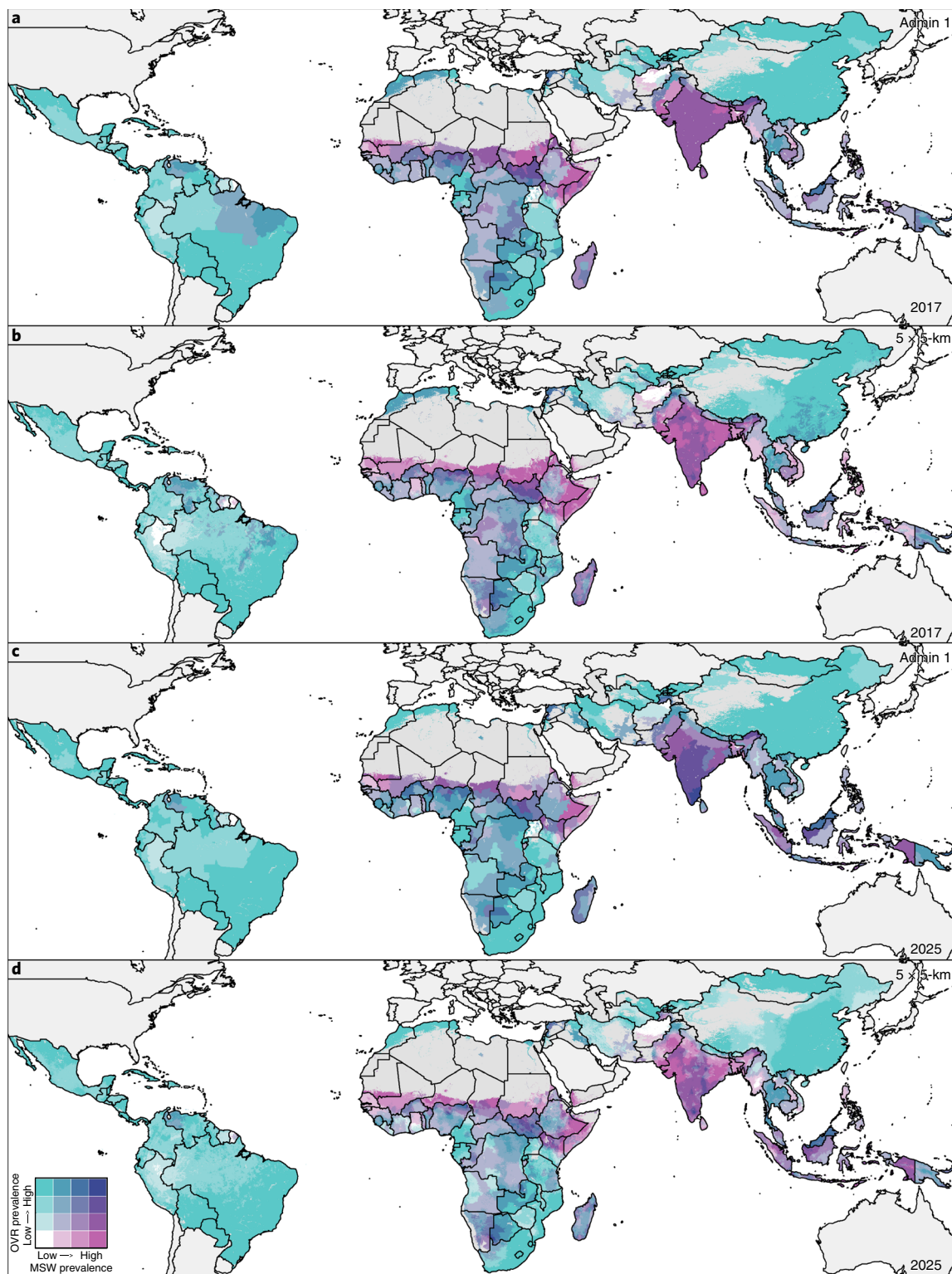
Across LMICs, trends in childhood overweight have increased while wasting decreased by different magnitudes from 2000–2017, leading to the emergence of DBM in several areas. As countries experience economic growth, they may undergo nutritional transitions wherein the challenges of undernutrition are replaced by those of overweight or the co-occurrence of both conditions<sup>4</sup>. Overall, food security has improved across LMICs in the past decade, which has led to increased availability of calories at the population level<sup>23</sup>. Although overweight is a reflection of excess calorie intake and reduced energy expenditure, there is a growing recognition that at the root of the rising rates of overweight are complex interactions between societal, environmental, food industry and individual factors, including biological, psychological and economical factors<sup>24</sup>. Understanding the factors underpinning these trends is key to predicting how nutrition programs can accelerate amelioration of wasting without incurring high rates of childhood overweight.

Although we included urbanicity as a covariate in our models, we were unable to reliably stratify our results by urban and rural areas. Urbanization is widely viewed as a key driver of the rise in overweight, but an increase in rural body mass index has recently been recognized as a main driver of the global epidemic of obesity in adults<sup>25</sup>. Such an analysis would thus add important context to our estimates. Case studies in China, Egypt, India, Mexico, the Philippines and South Africa have demonstrated a consistent trend of increased energy content of diets<sup>26</sup>. Relatively rural areas in China have experienced an increase in the intake of animal source foods and edible oils, likely due to the decreasing cost of these products. Further, increased use of motor vehicles and labor-saving technologies in agriculture have caused a decrease in energy expenditure in all these countries. In Brazil, household consumption of high-calorie ultra-processed foods has steadily replaced that of fresh or minimally processed foods<sup>27</sup>. Nutritious diets consisting of the latter can help prevent both wasting and stunting, thus work is needed to identify how dietary patterns differ between wasted and overweight children and the underlying factors causing those differences. Widespread collection and assembly of nutrition data from older children and adults would also contribute to a more complete understanding of longitudinal nutrition patterns.

In addition to tracking progress, child nutrition measurements are important for predicting and averting morbidity and mortality. Wasting is often indicative of short-term weight loss due to food shortages, famine or diseases such as diarrhea<sup>28–30</sup> and puts children at greater risk of succumbing to common infections<sup>28</sup>. Childhood overweight is likely to progress into adulthood and is associated with NCDs<sup>24</sup>, including cardiovascular disease, type 2 diabetes, sleep apnea and cancer<sup>31,32</sup>. Routine monitoring and reporting of child nutrition status can highlight trends and act as an early warning for health systems, particularly in the context of epidemiological transitions<sup>4</sup>.

Although overall spending on development assistance and investments to address malnutrition from government donors have remained steady, those from multilateral institutions have increased since 2013, amounting to US\$856 million in overseas development assistance in 2016 (ref. 15). These investments, however, fall short of the estimated US\$3.5 trillion per year that malnutrition costs society, US\$500 billion of which is attributable to overweight and obesity<sup>33</sup>. By focusing on prevention and early action, healthcare costs can be reduced and human capital increased. One difficulty, however, is addressing the different forms of malnutrition in tandem. Multiple forms of malnutrition are the new normal, according to the GNR<sup>15</sup> and Scaling Up Nutrition<sup>34,35</sup>. Double-duty actions that could simultaneously combat undernutrition, overweight, obesity, and diet-related NCDs have been proposed to address this problem<sup>36–38</sup>. Despite progress in identifying such actions, such as the promotion of breastfeeding, double-duty approaches have not been widely adopted. To better respond to the diverse and rapidly





**Fig. 5 | Overlapping population-weighted quartiles of overweight and wasting prevalence in children under 5 across LMICs in 2017 and 2025.**

**a–d**, Prevalence of moderate-to-severe overweight (OVR) and wasting (MSW) among children under 5 years of age in 2017 at the first administrative unit (**a**) and at a 5 × 5-km resolution (**b**). **c,d**, Estimated prevalence of moderate to severe OVR and MSW among children under 5 years of age in 2025 at the first administrative unit (**c**) and at a 5 × 5-km resolution (**d**). Quartile cutoffs were 0–5%, ≥5–10%, ≥10–15% and ≥15%. Maps reflect administrative boundaries, land cover, lakes and population; gray colored areas have grid cells classified as ‘barren or sparsely vegetated’ and had fewer than ten people per 1 × 1-km grid cell in 2017 or were not included in these analyses<sup>39–45</sup>. Maps were generated using ArcGIS Desktop 10.6.

evolving nutrition challenges facing LMICs, sustainable and health-promoting food systems are needed to slow the development of DBM. Due to the multiple causality of malnutrition, multisector

collaboration is required, including agriculture, trade and industry, environment, communication and education, all working towards policy and intervention coherence<sup>8,24</sup>.

There are several limitations to these analyses, mainly concerning the quantity and quality of the underlying data in the models, as shown in our uncertainty maps (Figs. 1f and 2f). Missing or improbable values in the primary data may contribute bias in the estimates and thus we have incorporated covariates to improve the estimates in areas where data are sparse. Additionally, differences in measurement techniques between surveys, scale miscalibration or equipment failure and poor training and standardization of measurers may contribute bias. Although our estimates were produced at a high spatial resolution, they were limited to prevalence by area, rather than the co-occurrence of wasting and overweight experienced by the same households or individuals. Additional work is required to identify the immediate and basic causes that lead to both wasting and obesity coexisting in the same geographical areas so that appropriate solutions can be identified. Future studies will consider maternal indicators associated with child nutritional outcomes, such as anemia and examine the co-distribution of overweight and stunting to broaden our assessment. New modeling approaches are currently in development to provide full distributions of height, weight and age, for more complete assessments of DBM using all important indicators of undernutrition.

Commendable gains have been made globally against child malnutrition over the past two decades. Our mapped estimates, however, show that high rates of wasting persist and overweight is increasing among young children in many LMICs. Identifying the causes underlying the presence of wasting or overweight in children living in the same community is necessary to formulate appropriate solutions. The estimates provided by this study can aid in the identification of specific areas where further insight can be gathered and trials of policy interventions administered, ultimately contributing to the UN Decade of Action on Nutrition process of sustained and coherent implementation of policies and programs<sup>37</sup>.

### Online content

Any methods, additional references, Nature Research reporting summaries, source data, extended data, supplementary information, acknowledgements, peer review information; details of author contributions and competing interests; and statements of data and code availability are available at <https://doi.org/10.1038/s41591-020-0807-6>.

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## Methods

**Overview.** Our study follows the Guidelines for Accurate and Transparent Health Estimates Reporting<sup>48</sup> (Supplementary Table 1). The analyses used model-based geostatistics to generate local-, administrative- and national-level estimates of children under 5 overweight, wasting prevalence and double burden in LMICs over time. Using an ensemble modeling framework that fed into a Bayesian generalized linear mixed-effects model with a correlated space-time random effect and 1,000 draws from an approximate posterior distribution, we generated annual prevalence estimates for overweight and wasting on a 5 × 5-km grid over 105 LMICs from 2000 to 2017 and aggregated these to administrative and national levels (Supplementary Table 2). Countries were selected for inclusion in this study using the socio-demographic index (SDI), a summary measure of development that combines education, fertility and poverty<sup>47</sup>. Selected countries were in the low, lower-middle and middle SDI quintiles, with several exceptions (Supplementary Table 2). China, Libya, Malaysia, Panama and Turkmenistan were included despite higher-middle SDIs for geographic continuity with other included countries. Albania, Bosnia-Herzegovina and Moldova were excluded due to geographic discontinuity and lack of available survey data. We did not conduct estimates for the island nations of American Samoa, Federated States of Micronesia, Fiji, Kiribati, Marshall Islands, Samoa, Solomon Islands or Tonga, as no survey data could be sourced.

**Data. Surveys and child anthropometry data.** We extracted individual-level height, weight and age data for children under 5 from household survey series including the Demographic and Health Surveys, Multiple Indicator Cluster Surveys, Living Standards Measurement Study and Core Welfare Indicators Questionnaire, among other country-specific child health and nutrition surveys<sup>49–52</sup> (Supplementary Tables 3 and 4). Included in our models were 420 georeferenced household surveys representing over 3 million children under 5. Each individual child record was associated with a cluster, a group of neighboring households or a ‘village’ that acted as a primary sampling unit. Approximately 185 surveys with height, weight and age data included geographic coordinates or precise place names for each cluster within that survey. In the absence of geographic coordinates for each cluster, we assigned data to the smallest available administrative area unit in the survey (polygon) while accounting for the survey sample design (15,781 survey polygons for overweight and wasting)<sup>53,54</sup>. Boundary information for these administrative units was obtained as shapefiles either directly from the surveys or by matching to shapefiles in the Global Administrative Unit Layers<sup>55</sup> database or the Database of Global Administrative Areas<sup>56</sup>. In select cases, shapefiles provided by the survey administrator were used or custom shapefiles were created based on survey documentation. These area data were resampled to point locations using a population-weighted sampling approach over the relevant area unit with the number of locations set proportionally to the number of grid cells in the area and the total weights of all the resampled points summing to one<sup>45</sup>.

Select data sources were excluded for the following reasons: missing survey weights for areal data, missing sex or age variable, incomplete sampling (for example, only children ages 0–3 years measured) or untrustworthy data (as determined by the survey administrator or by inspection). Details on the survey data excluded for each country can be found in Supplementary Table 5. Data extraction and processing methods have been described previously<sup>21</sup>.

**Child anthropometry.** Using height, weight, age and sex data for each individual, WHZs were calculated using the age-, sex- and indicator-specific lambda-mu-sigma values from the 2006 WHO Child Growth Standards<sup>10,57</sup>. The lambda-mu-sigma methodology allows for Gaussian *z* score calculations and comparisons to be applied to skewed, non-Gaussian distributions<sup>58</sup>. A child was classified as overweight or wasted if their weight-for-height/length was more than two s.d. (*z* scores) above or below the WHO growth reference population, respectively<sup>59</sup>. These individual-level data observations were then collapsed to cluster-level totals for the number of children sampled and total number of children under 5 affected by overweight and the total number of children who are wasted out of the children who were not overweight.

**Temporal resolution.** We estimated prevalence of overweight and wasting annually from 2000 to 2017 using a model that allowed us to account for data points measured across survey years, and as such, allows us to predict at monthly or finer temporal resolutions. We were limited, however, both computationally and by the temporal resolution of covariates (Supplementary Table 6) and thus produced only annual estimates.

**Seasonality adjustment.** WHZs were used to calculate individual child wasting status. As a data preprocessing step, we performed a seasonality adjustment on individual-level child weights in order to account for differences in observed child weight that may have been due to food scarcity during the month in which the survey was conducted. To adjust weight measurements, we fitted a model for each region with a 12-month seasonal spline, a country-level fixed effect and a smooth spline over the duration of our data collection using the *mgcv* package in R and the following formula:

$$\text{WHZ} \sim s_{cc}(\text{month}) + s_p(t) + \text{as.factor}(\text{country}).$$

Month is the integer-valued month of the year (1, ..., 12), *t* is the time of the interview in integer months since the earliest observation of any child in the dataset and country is a factor variable representing the country where the observation was recorded. We modeled the periodic component on months using 12 cyclic cubic (*cc*) regression splines basis functions and we accounted for a smooth longer time temporal trend using four thin-plate (*tp*) splines. The country effects and the long-term temporal spline were included only to avoid confounding during fitting of the seasonal spline fit and neither country effects nor the long-term trend was used in the seasonal adjustment. We then adjusted all observations to account for the difference in the seasonal period between the month of the interview and an average day of the year as determined by which days aligned with the mean of the periodic spline.

**Spatial covariates.** In order to leverage strength from locations with observations to the entire spatial-temporal domain, we compiled several 5 × 5-km raster layers of putative socioeconomic and environmental correlates of malnutrition in the 105 LMICs (Supplementary Table 6). These covariates were selected based on their potential to be predictive for overweight and wasting, according to literature review and plausible hypothesis as to their influence. Acquisition of temporally dynamic datasets, where possible, was prioritized to best match our observations and thus predict the changing dynamics of the two indicators. Of the 12 covariates included, 6 were temporally dynamic and were reformatted as a synoptic mean over each estimation period or as a mid-period year estimate. These included average daily mean rainfall (precipitation), educational attainment in women of reproductive age (15–49 years old), enhanced vegetation index, fertility, urbanicity and population. The remaining six covariate layers were static throughout the study period and were applied uniformly across all modeling years; these covariates included growing season length, irrigation, nutritional yield for vitamin A, nutritional yield for protein, nutritional yield for iron and travel time to nearest settlement >50,000 inhabitants.

To select covariates and capture possible nonlinear effects and complex interactions between them, an ensemble covariate modeling method was implemented<sup>60</sup>. For each region, three submodels were fitted to our dataset, using all of our covariate data as explanatory predictors: generalized additive models, boosted regression trees and lasso regression. Each submodel was fitted using fivefold cross-validation to avoid overfitting and the out-of-sample predictions from across the five holdouts were compiled into a single comprehensive set of out-of-sample predictions from that model. Additionally, the same submodels were also run using 100% of the data and a full set of in-sample predictions were created. The three sets of out-of-sample submodel predictions were fed into the full geostatistical model as the explanatory covariates when performing the model fitting. The in-sample predictions from the submodels were used as covariates when generating predictions using the fitted full geostatistical model. A recent study has shown that this ensemble approach can improve predictive validity by up to 25% over an individual model<sup>60</sup>.

**Analysis. Geostatistical model.** In this study, wasting was defined as the proportion of children under 5 below negative 2 WHZ (<−2 WHZ); normal category, the proportion of children under 5 between negative 2 and positive 2 WHZ *z* score (>−2 and <2 WHZ); and overweight was defined as the proportion of children under 5 above positive 2 WHZ *z* score (>2 WHZ) as defined in the WHO growth reference population<sup>59</sup>. To model the full distribution of possible indicators of nutritional status in WHZ (wasting (<−2 WHZ), normal (>−2 and <2 WHZ) and overweight (>2 WHZ)), we used an ordinal modeling approach<sup>61,62</sup> to estimate the relative proportion of each indicator. A similar modeling approach was used to estimate vaccine coverage in Africa<sup>63</sup>.

We used a continuation ratio model to estimate the prevalence of three categories: wasting, normal weight and overweight. We first modeled the proportion of wasting children within a Bayesian hierarchical framework using logistic regression with a spatially and temporally explicit generalized linear mixed-effects model. Second, we modeled the proportion of the children that were overweight conditioned on not being wasted using the same Bayesian modeling framework. The estimates from the second conditional model were then combined with the wasting estimates to compute the proportion of overweight children in the full distribution.

At each cluster, *j*, where *j* = 1, 2, ..., *n*, and time *t*, where *t* = 2000, 2001, ..., 2017, the prevalence of wasting was modeled using the observed number of children in cluster *d*, who were found to be wasted as a binomial count data *C<sub>d</sub>* among a sample size *N<sub>d</sub>*.

$$\begin{aligned} C_d | p_{i(d),t(d)}, N_d &\sim \text{Binomial}(p_{i(d),t(d)}, N_d) \forall \text{ observe clusters } d \text{ logit}(p_{i,t}) \\ &= \beta_0 + \mathbf{X}_{i,t} \boldsymbol{\beta} + Z_{i,t} + \epsilon_{ctr(i)} + \epsilon_{i,t} + Z_{i,t} \forall i \in \text{spatial domain} \forall t \in \text{time domain} \\ \sum_{h=1}^3 \beta_h &= 1 \\ \epsilon_{ctr} &\sim \text{iid Normal}(0, \gamma^2) \\ \epsilon_{i,t} &\sim \text{iid Normal}(0, \sigma^2) \\ \mathbf{Z} &\sim \text{GP}(0, \Sigma^{\text{space}} \otimes \Sigma^{\text{time}}) \\ \Sigma^{\text{space}} &= \frac{\omega^2}{1 + (\nu)^{2\alpha-1}} \times (\kappa D)^{\nu} \times \mathbf{K}_{\nu}(\kappa D) \\ \Sigma_{j,k}^{\text{time}} &= \rho^{|k-j|} \end{aligned}$$



For indices  $d$ ,  $i$  and  $t$ ,  $*$ (index) is the value of  $*$  at the index. The annual prevalence of wasting,  $p_{i,t}$ , in cluster  $i$ , in time  $t$ , was modeled as a linear combination of the three submodels, (generalized additive models, boosted regression trees and lasso regression), rasterized covariate values,  $X_{i,t}$ , a correlated spatiotemporal random effect term  $Z_{i,t}$ , country random effects  $\epsilon_{ctr(i)}$ , with one unstructured country random effect fitted for each country in the modeling region and all  $\epsilon_{ctr}$  sharing a common variance parameter,  $\gamma^2$ , and an independent nugget random effect  $\epsilon_{i,t}$ , with variance parameter,  $\sigma^2$ . Coefficients  $\beta_h$  in the three submodels  $h=1, 2, 3$  represent their respective predictive weighting in the logit link, while the joint error term  $Z_{i,t}$  accounts for residual spatiotemporal autocorrelation between individual data points that remain after accounting for the predictive effect of the submodel covariates, the country-level random effect  $\epsilon_{ctr(i)}$  and the nugget,  $\epsilon_{i,t}$ . The residuals  $Z_{i,t}$  were modeled as a three-dimensional Gaussian process in space–time centered at zero and with a covariance matrix constructed from a Kronecker product of spatial and temporal covariance kernels. The spatial covariance,  $\Sigma^{\text{space}}$ , was modeled using an isotropic and stationary Matérn function<sup>64</sup> and temporal covariance,  $\Sigma^{\text{time}}$ , as an annual autoregressive (AR1) function over the 18 years represented in the model. In the stationary Matérn function,  $\Gamma$  is the gamma function,  $K_\nu$  is the modified Bessel function of order  $\nu > 0$ ,  $\kappa > 0$  is a scaling parameter,  $D$  denotes the Euclidean distance and  $\omega^2$  is the marginal variance. The scaling parameter,  $\kappa$ , is defined to be  $\kappa = \sqrt{8\nu}/\delta$ , where  $\delta$  is a range parameter (about the distance where the covariance function approaches 0.1) and  $\nu$  is a scaling constant, which is set to 2 rather than fitted from the data. The number of rows and the number of columns of the spatial Matérn covariance matrix are both equal to the number of spatial mesh points for a given modeling region. The number of rows and the number of columns of the spatial Matérn covariance matrix are both equal to the number of spatial mesh points for a given modeling region. In the AR1 function,  $\rho$  is the autocorrelation function and  $k$  and  $j$  are points in the time series where  $|k-j|$  defines the lag. The number of rows and the number of columns of the AR1 covariance matrix are both equal to the number of temporal mesh points (18). The number of rows and the number of columns of the space–time covariance matrix,  $\Sigma^{\text{space}} \otimes \Sigma^{\text{time}}$ , for a given modeling region are both equal to the number of spatial mesh points  $\times$  the number of temporal mesh points.

This approach leverages the residual correlation structure to more accurately predict prevalence estimates for locations with no data, while also propagating the dependence in the data through to uncertainty estimates<sup>65</sup>. The posterior distributions were fitted using computationally efficient and accurate approximations in R-INLA<sup>66,67</sup> (integrated nested Laplace approximation) with the stochastic partial differential equations<sup>68</sup> approximation to the Gaussian process residuals using R project v.3.5.1. The stochastic partial differential equations approach using INLA has been demonstrated elsewhere, including the estimation of health indicators, particulate air matter and population age structure<sup>69–71</sup>. Uncertainty intervals were generated from 1,000 draws (statistically plausible candidate maps)<sup>72</sup> created from the posterior-estimated distributions of modeled parameters.

**Post estimation.** To transform grid cell-level estimates into a range of information useful to a wide constituency of potential users, estimates were aggregated at first and second administrative units specific to each country and at national levels<sup>73</sup>. Although the models can predict all locations covered by available raster covariates, all final model outputs for which land cover was classified as ‘barren or sparsely vegetated’ on the basis of Moderate Resolution Imaging Spectroradiometer (MODIS) satellite data (2013) were masked<sup>74</sup>. Areas where the total population density was less than ten individuals per  $1 \times 1$ -km grid cell in 2015 were also masked in the final outputs.

**Model validation.** Models were validated using spatially stratified fivefold out-of-sample cross-validation. In order to offer a more stringent analysis by accounting for some of the spatial correlation in the data, holdout folds were created by combining sets of all data falling with first administrative level areas. Validation was performed by calculating bias (mean error), variance (root-mean-square error), 95% data coverage within prediction intervals and correlation between observed data and predictions. All validation metrics were calculated on the out-of-sample predictions from the fivefold cross-validation. All validation procedures and corresponding results are provided in Supplementary Tables 7–18.

**Projections.** To compare our estimated rates of improvement in overweight and wasting prevalence over the last 18 years with the improvements needed between 2017 and 2025 to meet WHO GNTs, we performed a simple projection using estimated AROC applied to the final year of our estimates. Both AROC and projections were calculated at the draw-level to obtain the uncertainty of the estimates.

For each indicator  $i$ , we calculated AROC at each grid cell ( $m$ ) by calculating the AROC between each pair of adjacent years  $t$ :

$$\text{AROC}_{u,m,t} = \text{logit} \left( \frac{p_{u,m,t}}{p_{u,m,t-1}} \right)$$

We then calculated a weighted AROC for each indicator by taking a weighted average across the years, where more recent AROCs were given more weight in the average. We defined the weights to be:

$$W_t = (t - 2000 + 1)^\gamma,$$

where  $\gamma$  may be chosen to give varying amounts of weight across the years. For each indicator, we then calculated the average AROC to be:

$$\text{AROC}_{u,m} = \left( \sum_{2001}^{2017} W_t \times \text{AROC}_{u,m,t} \right)$$

Finally, we calculated the projections (Proj) by applying the AROC in our 2017 mean prevalence estimates to produce estimates in 8 years from 2017 to 2025.

$$\text{Proj}_{u,m,2025} = \text{logit}^{-1} \left( \text{logit}(p_{u,m,2017}) + \text{AROC}_{u,m} \times 8 \right).$$

This projection scheme is analogous to the methods used in the Global Burden of Disease 2017 study<sup>47</sup> for measurement of progress and projected attainment of health-related Sustainable Development Goals. Our projections are based on the assumption that areas will sustain the current AROC, and the precision of the AROC estimates is dependent on the level of uncertainty emanating from the estimation of annual prevalence.

**Priors.** The following priors were used for our overweight and wasting models:

$$\begin{aligned} \beta_0 &\sim N(\mu = 0, \sigma^2 = 3^2), \\ \beta &\sim \text{iid } N\left(\mu = \frac{1}{\text{no. ensemble models}}, \sigma^2 = 3^2\right), \\ \log\left(\frac{1+\rho}{1-\rho}\right) &\sim N(\mu = 4, \sigma^2 = 1.2^2), \\ \log\left(\frac{1}{\sigma_{\text{nugget}}^2}\right) &\sim \text{loggamma}(\alpha = 1, \gamma = 5 \times 10^{-5}), \\ \log\left(\frac{1}{\sigma_{\text{country}}^2}\right) &\sim \text{loggamma}(\alpha = 1, \gamma = 5 \times 10^{-5}), \\ \theta_1 = \log(\sigma_k^2) &\sim N(\mu_{\theta_1}, \sigma_{\theta_1}^2) \\ \theta_2 = \log(\kappa) &\sim N(\mu_{\theta_2}, \sigma_{\theta_2}^2) \end{aligned}$$

Given that our covariates used in INLA (the predicted outputs from the ensemble models) should be on the same scale as our predictive target, we believe that the intercept in our model should be close to zero and that the regression coefficients should sum to 1. As such, we chose the prior for our intercept to be  $N(0, \sigma^2 = 3^2)$  and the prior for the fixed-effect coefficients to be  $N\left(\frac{1}{\text{no. ensemble models}}, \sigma^2 = 3^2\right)$ . The prior on the temporal correlation parameter,  $\rho$ , was chosen to be mean zero, showing no prior preference for either positive or negative autocorrelation structure and with a distribution wide enough such that within three s.d. of the mean, the prior includes values of  $\rho$  ranging from  $-0.95$  to  $0.95$ . The priors on the random effect variances were chosen to be relatively loose given that we believe our fixed-effects covariates should be well correlated with our outcome of interest, which might suggest relatively small random effects values. At the same time, we wanted to avoid using a prior that was so diffuse as to actually put high prior weight on large random effect variances. For stability, we used the uncorrelated multivariate normal priors that INLA automatically determines (based on the finite elements mesh) for the log-transformed spatial hyperparameters  $\kappa$  and  $\tau$ . In our parameterization, we represent  $\alpha$  and  $\gamma$  in the log gamma distribution as shape and inverse-scale, respectively.

**Prior sensitivity analysis.** Sensitivity analysis was undertaken to assess the impact of the hyper-priors for the nugget, country random effects, and space–time correlation. We considered two different sets of priors related to the nugget and country random effects and three set related to space–time correlation, resulting in six different combinations of hyper-priors as outlined below.

**Model 1:** In this model, we used the default hyper-priors in INLA<sup>75</sup> (for both nugget and country random effects. The hyper-prior for the AR1 rho,  $\rho$ , was retained as shown below.

$$\begin{aligned} \log\left(\frac{1}{\sigma_{\text{nugget}}^2}\right) &\sim \text{loggamma}(\alpha = 1, \gamma = 5 \times 10^{-5}) \text{ and} \\ \log\left(\frac{1}{\sigma_{\text{country}}^2}\right) &\sim \text{loggamma}(\alpha = 1, \gamma = 5 \times 10^{-5}). \\ \log\left(\frac{1+\rho}{1-\rho}\right) &\sim \text{Normal}(\mu = 4, \sigma^2 = 1.2^2) \end{aligned}$$

**Model 2:** The hyper-priors for nugget were changed as indicated below, where hyper-priors for country random effect were the default hyper-priors in INLA. The hyper-priors for the AR1 rho,  $\rho$ , were retained the same as model 1.

$$\begin{aligned} \log\left(\frac{1}{\sigma_{\text{nugget}}^2}\right) &\sim \text{loggamma}(\alpha = 1, \gamma = 2) \text{ and} \\ \log\left(\frac{1}{\sigma_{\text{country}}^2}\right) &\sim \text{loggamma}(\alpha = 1, \gamma = 5 \times 10^{-5}). \\ \log\left(\frac{1+\rho}{1-\rho}\right) &\sim \text{Normal}(\mu = 4, \sigma^2 = 1.2^2) \end{aligned}$$

Model 3: In this model the hyper-priors for country random effects and nugget were exchanged, where hyper-priors for nugget were the default hyper-priors in INLA. The hyper-priors for the AR1 rho,  $\rho$ , were retained the same as model 1.

$$\begin{aligned}\log\left(\frac{1}{\sigma_{\text{nug}}^2}\right) &\sim \text{loggamma}(\alpha = 1, \gamma = 5 \times 10^{-5}) \text{ and} \\ \log\left(\frac{1}{\sigma_{\text{country}}^2}\right) &\sim \text{loggamma}(\alpha = 1, \gamma = 2). \\ \log\left(\frac{1+\rho}{1-\rho}\right) &\sim \text{Normal}(\mu = 4, \sigma^2 = 1.2^2)\end{aligned}$$

Model 4: In this model, we used the default hyper-priors in INLA for less informative nugget and country random effects. The hyper-priors for the AR1 rho,  $\rho$ , were changed.

$$\begin{aligned}\log\left(\frac{1}{\sigma_{\text{nugget}}^2}\right) &\sim \text{loggamma}(\alpha = 1, \gamma = 5 \times 10^{-5}) \text{ and} \\ \log\left(\frac{1}{\sigma_{\text{country}}^2}\right) &\sim \text{loggamma}(\alpha = 1, \gamma = 5 \times 10^{-5}). \\ \log\left(\frac{1+\rho}{1-\rho}\right) &\sim \text{Normal}(\mu = 0, \sigma^2 = 1.2^2)\end{aligned}$$

Model 5: In this model, we used the default hyper-priors in INLA for both nugget and country random effects. The hyper-priors for the AR1 rho,  $\rho$ , were the default in INLA.

$$\begin{aligned}\log\left(\frac{1}{\sigma_{\text{nugget}}^2}\right) &\sim \text{loggamma}(\alpha = 1, \gamma = 5 \times 10^{-5}) \text{ and} \\ \log\left(\frac{1}{\sigma_{\text{country}}^2}\right) &\sim \text{loggamma}(\alpha = 1, \gamma = 5 \times 10^{-5}). \\ \log\left(\frac{1+\rho}{1-\rho}\right) &\sim \text{Normal}(\mu = 0, \sigma^2 = 2.58^2)\end{aligned}$$

The predicted estimates for all models with different sets of hyper-priors were highly correlated at the grid-cell level and yielded low mean absolute differences (Supplementary Table 7). We ultimately selected the less informative priors for nugget and country random effects as they are default priors in the INLA package and have been applied widely<sup>76,77</sup> and selected a more stringent parameterization of our space-time correlation, as indicated in model 1.

**Mesh construction.** We constructed the finite elements mesh for the stochastic partial differential equation approximation to the Gaussian process regression using a simplified polygon boundary (in which coastlines and complex boundaries were smoothed) for each of the regions within our model. We set the inner mesh triangle maximum edge length (the mesh size for areas over land) to be 0.75 degrees and the buffer maximum edge length (the mesh size for areas over the ocean) to be 5 degrees. An example finite elements mesh constructed for Eastern SSA mesh is described by Kinyoki et al.<sup>21</sup>.

**Reporting Summary.** Further information on research design is available in the Nature Research Reporting Summary linked to this article.

## Data availability

Our study follows the Guidelines for Accurate and Transparent Health Estimates Reporting<sup>48</sup> (Supplementary Table 1). The findings of this study are supported by data available in public online repositories, data publicly available upon request of the data provider and data not publicly available due to restrictions by the data provider. Nonpublicly available data were used under license for the current study but may be available from the authors upon reasonable request and with permission of the data provider. Details of data sources and availability can be found in Supplementary Tables 2–5. The full output of the analyses are publicly available in the Global Health Data Exchange (<http://ghdx.healthdata.org/record/ihme-data/lmic-double-burden-of-malnutrition-geospatial-estimates-2000-2017>) and can further be explored via customized data visualization tools (<https://vizhub.healthdata.org/lbd/dbm>). Administrative boundaries were retrieved from the Database of Global Administrative Areas<sup>39</sup>. Land cover was retrieved from the online Data Pool, courtesy of the NASA EOSDIS Land Processes Distributed Active Archive Center, USGS/Earth Resources Observation and Science Center, Sioux Falls, South Dakota<sup>40</sup>. Lakes were retrieved from the Global Lakes and Wetlands Database, courtesy of the World Wildlife Fund and the Center for Environmental Systems Research, University of Kassel<sup>41,42</sup>. Populations were retrieved from WorldPop<sup>43,44</sup>.

## Code availability

All code used for these analyses is publicly available online at <http://ghdx.healthdata.org/record/ihme-data/lmic-double-burden-of-malnutrition-geospatial-estimates-2000-2017> and at <http://github.com/ihmeuw/lbd/tree/dbm-lmic-2020>.

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## Author contributions

D.K.K., J.M.R., A.A. and S.I.H. conceived and planned the study. A.L.-A. and D.K.K. obtained, extracted, processed and geopositioned data. D.K.K. carried out statistical analyses. The first draft of the manuscript was written by D.K.K., J.M.R., S.B.M., L.E.S., A.A. and S.I.H.; D.K.K., S.B.M. and J.M.R. finalized the manuscript based on comments from other authors and reviewer feedback. D.K.K., A.L.-A. and S.B.M. managed the Supplementary Information. All authors provided intellectual input into aspects of this study. Additional details on author contributions are in the Supplementary Information.

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## Additional information

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**Correspondence and requests for materials** should be addressed to S.I.H.

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## LBD Double Burden of Malnutrition Collaborators

Damaris K. Kinyoki<sup>1,2</sup>, Jennifer M. Ross<sup>1,3,4</sup>, Alice Lazzar-Atwood<sup>1</sup>, Sandra B. Munro<sup>1</sup>, Lauren E. Schaeffer<sup>1</sup>, Mahdiah Abbasalizad-Farhangi<sup>5</sup>, Masoumeh Abbasi<sup>6</sup>, Hedayat Abbastabar<sup>7</sup>, Ahmed Abdelalim<sup>8</sup>, Amir Abdoli<sup>9</sup>, Mohammad Abdollahi<sup>10</sup>, Ibrahim Abdollahpour<sup>11</sup>, Rizwan Suliankatchi Abdulkader<sup>12</sup>, Nebiyu Dereje Abebe<sup>13,14</sup>, Teshome Abuka Abebo<sup>15</sup>, Kedir Hussein Abegaz<sup>16,17</sup>, Hassan Abolhassani<sup>18,19</sup>, Lucas Guimarães Abreu<sup>20</sup>, Michael R. M. Abrigo<sup>21</sup>, Abdelrahman I. Abushouk<sup>22</sup>, Manfred Mario Kokou Accrombessi<sup>23</sup>, Dilaram Acharya<sup>24,25</sup>, Maryam Adabi<sup>26</sup>, Akindele Olupelumi Adebisi<sup>27,28</sup>, Isaac Akinkunmi Adedeji<sup>29</sup>, Victor Adekanmbi<sup>30</sup>, Abiodun Moshood Adeoye<sup>31,32</sup>, Olatunji O. Adetokunboh<sup>33,34</sup>, Davoud Adham<sup>35</sup>, Posi Emmanuel Aduroja<sup>36</sup>, Shailesh M. Advani<sup>37,38</sup>, Mohsen Afarideh<sup>39</sup>, Mohammad Aghaali<sup>40</sup>, Anurag Agrawal<sup>41,42</sup>, Tauseef Ahmad<sup>43,44</sup>, Keivan Ahmadi<sup>45</sup>, Sepideh Ahmadi<sup>46</sup>, Muktar Beshir Ahmed<sup>47</sup>, Rushdia Ahmed<sup>48,49</sup>, Olufemi Ajumobi<sup>50,51</sup>, Chalachew Genet Akal<sup>52</sup>, Temesgen Yihunie Akalu<sup>53</sup>, Tomi Akinyemiju<sup>54,55</sup>, Blessing Akombi<sup>56</sup>, Ziyad Al-Aly<sup>57,58</sup>, Samiah Alam<sup>59</sup>, Genet Melak Alamene<sup>60</sup>, Turki M. Alanzi<sup>61</sup>, Jacqueline Elizabeth Alcalde Rabanal<sup>62</sup>, Niguse Meles Alema<sup>63</sup>, Beriwan Abdulqadir Ali<sup>64,65</sup>, Muhammad Ali<sup>66</sup>, Mehran Alijanzadeh<sup>67</sup>, Cyrus Alinia<sup>68</sup>, Vahid Alipour<sup>69,70</sup>, Hesam Alizade<sup>71,72</sup>, Syed Mohamed Aljunid<sup>73,74</sup>, Afshin Almasi<sup>75</sup>, Amir Almasi-Hashiani<sup>76</sup>, Hesham M. Al-Mekhlafi<sup>77,78</sup>, Rajaa M. Al-Raddadi<sup>79</sup>, Khalid Altirkawi<sup>80</sup>, Nelson Alvis-Guzman<sup>81,82</sup>, Nelson J. Alvis-Zakzuk<sup>83,84</sup>, Azmeraw T. Amare<sup>85,86</sup>, Adeladza Kofi Amegah<sup>87</sup>, Saeed Amini<sup>88</sup>, Mostafa Amini Rarani<sup>89</sup>, Fatemeh Amiri<sup>90</sup>, Arianna Maeve Loreche Amit<sup>91,92</sup>, Nahla Hamed Anber<sup>93</sup>, Catalina Liliana Andrei<sup>94</sup>, Fereshteh Ansari<sup>95,96</sup>, Alireza Ansari-Moghaddam<sup>97</sup>, Zelalem Alamrew Anteneh<sup>98</sup>, Carl Abelardo T. Antonio<sup>99,100</sup>, Ernoiz Antriyandarti<sup>101</sup>, Davood Anvari<sup>102,103</sup>, Raziq Anwer<sup>104</sup>,

Seth Christopher Yaw Appiah<sup>105,106</sup>, Jalal Arabloo<sup>69</sup>, Morteza Arab-Zozani<sup>107</sup>, Ephrem Mebrahtu Araya<sup>63</sup>, Zohreh Arefi<sup>108</sup>, Olatunde Aremu<sup>109</sup>, Johan Ärnlov<sup>110,111</sup>, Afsaneh Arzani<sup>112,113</sup>, Mehran Asadi-Aliabadi<sup>114</sup>, Ali A. Asadi-Pooya<sup>115</sup>, Samaneh Asgari<sup>116</sup>, Babak Asghari<sup>117</sup>, Alebachew Fasil Ashagre<sup>118</sup>, Anemaw A. Asrat<sup>98</sup>, Bahar Ataeinia<sup>119</sup>, Hagos Tasew Atalay<sup>120</sup>, Desta Debalkie Atnafu<sup>121</sup>, Maha Moh'd Wahbi Atout<sup>122</sup>, Marcel Ausloos<sup>123,124</sup>, Euripide F. G. A. Avokpaho<sup>125,126</sup>, Ashish Awasthi<sup>127</sup>, Beatriz Paulina Ayala Quintanilla<sup>128,129</sup>, Martin Amogre Ayanore<sup>130</sup>, Yared A. Asmare Aynalem<sup>131</sup>, Abbas Azadmehr<sup>132</sup>, Samad Azari<sup>69</sup>, Ghasem Azarian<sup>133</sup>, Zelalem Nigussie Azene<sup>134</sup>, Ebrahim Babaee<sup>114</sup>, Alaa Badawi<sup>135,136</sup>, Ashish D. Badiye<sup>137</sup>, Mohamad Amin Bahrami<sup>138</sup>, Atif Amin A. Baig<sup>139,140</sup>, Ahad Bakhtiari<sup>141</sup>, Shankar M. Bakkannavar<sup>142</sup>, Senthilkumar Balakrishnan<sup>143</sup>, Ayele Geleto Bali<sup>144</sup>, Maciej Banach<sup>145,146</sup>, Palash Chandra Banik<sup>147</sup>, Zahra Baradaran-Seyed<sup>148</sup>, Adhanom Gebreegziabher Baraki<sup>53</sup>, Miguel A. Barboza<sup>149,150</sup>, Till Winfried Bärnighausen<sup>151,152</sup>, Lingkan Barua<sup>147</sup>, Huda Basaleem<sup>153</sup>, Sanjay Basu<sup>154</sup>, Mohsen Bayati<sup>155</sup>, Mulat Tirfie Bayih<sup>156</sup>, Habtamu Wondifraw Baynes<sup>157</sup>, Neeraj Bedi<sup>158,159</sup>, Masoud Behzadifar<sup>160</sup>, Meysam Behzadifar<sup>161</sup>, Yibeltal Alemu Bekele<sup>162</sup>, Derrick A. Bennett<sup>163</sup>, Dessalegn Ajema Berbada<sup>164</sup>, Kidanemariam Berhe<sup>165</sup>, Abadi Kidanemariam Berhe<sup>166,167</sup>, Adam E. Berman<sup>168</sup>, Robert S. Bernstein<sup>169,170</sup>, Reshmi Bhageerathy<sup>171</sup>, Dinesh Bhandari<sup>172,173</sup>, Pankaj Bharadwaj<sup>174,175</sup>, Natalia V. Bhattacharjee<sup>1</sup>, Kritika Bhattacharyya<sup>176,177</sup>, Ali Bijani<sup>178</sup>, Boris Bikbov<sup>179</sup>, Ver Bilano<sup>180</sup>, Nigus Bililign<sup>181</sup>, Muhammad Shahdaat Bin Sayeed<sup>182,183</sup>, Setognal Birara<sup>184</sup>, Minuye Biniam Biniam Birhane<sup>185</sup>, Minyichil Birhanu<sup>186</sup>, Raaj Kishore Biswas<sup>187,188</sup>, Zebeay Workneh Bitew<sup>189</sup>, Kassawmar Angaw Bogale<sup>98</sup>, Somayeh Bohloul<sup>190</sup>, Srinivasa Rao Bolla<sup>191</sup>, Archith Boloor<sup>192</sup>, Antonio M. Borzi<sup>193</sup>, Shiva Borzouei<sup>194</sup>, Oliver J. Brady<sup>195</sup>, Nicola Luigi Bragazzi<sup>196</sup>, Dejana Braithwaite<sup>197</sup>, Nikolay Ivanovich Briko<sup>198</sup>, Gabrielle Britton<sup>199</sup>, Shyam S. Budhathoki<sup>200</sup>, Sharath Burugina Nagaraja<sup>201</sup>, Reinhard Busse<sup>202</sup>, Zahid A. Butt<sup>203,204</sup>, Lucero Cahuana-Hurtado<sup>62</sup>, Luis Alberto Cámara<sup>205,206</sup>, Ismael R. Campos-Nonato<sup>207</sup>, Jorge Cano<sup>208</sup>, Josip Car<sup>209,210</sup>, Rosario Cárdenas<sup>211</sup>, Juan J. Carrero<sup>212</sup>, Félix Carvalho<sup>213</sup>, João Mauricio Castaldelli-Maia<sup>214</sup>, Carlos A. Castañeda-Orjuela<sup>215,216</sup>, Franz Castro<sup>199</sup>, Ester Cerin<sup>217,218</sup>, Collins Chansa<sup>219,220</sup>, Jaykaran Charan<sup>221</sup>, Pranab Chatterjee<sup>222</sup>, Vijay Kumar Chattu<sup>223</sup>, Bal Govind Chauhan<sup>224,225</sup>, Ali Reza Chavshin<sup>226</sup>, Mohammad Chehraz<sup>227,228</sup>, Tesfaye Yitna Chichiabellu<sup>229</sup>, Ken Lee Chin<sup>230</sup>, Devasahayam J. Christopher<sup>231</sup>, Dinh-Toi Chu<sup>232</sup>, Flavia M. Cicuttini<sup>233</sup>, Michael L. Collison<sup>1</sup>, Michael A. Cork<sup>1</sup>, Natalie Cormier<sup>1</sup>, Paolo Angelo Cortesi<sup>234</sup>, Vera M. Costa<sup>213</sup>, Abel Fekadu Fekadu Dadi<sup>235,236</sup>, Baye Dagnew<sup>237</sup>, Saad M. A. Dahlawi<sup>238</sup>, Giovanni Damiani<sup>239,240</sup>, Amira Hamed Darwish<sup>241</sup>, Ahmad Daryani<sup>242</sup>, Jai K. Das<sup>243</sup>, Rajat Das Gupta<sup>48,244</sup>, Claudio Dávila-Cervantes<sup>245</sup>, Nicole Davis Weaver<sup>1</sup>, Diego De Leo<sup>246</sup>, Jan-Walter De Neve<sup>151</sup>, Feleke Mekonnen Demeke<sup>52</sup>, Asmamaw Bizuneh Demis<sup>247,248</sup>, Dereje Bayissa Demissie<sup>249,250</sup>, Gebre Teklemariam Demoz<sup>251,252</sup>, Edgar Denova-Gutiérrez<sup>253</sup>, Kebede Deribe<sup>13,254</sup>, Rupak Desai<sup>255</sup>, Beruk Berhanu Desalegn<sup>256</sup>, Assefa Desalew<sup>257</sup>, Aniruddha Deshpande<sup>1</sup>, Sagnik Dey<sup>258</sup>, Samath Dhamminda Dharmaratne<sup>1,259</sup>, Preeti Dhillon<sup>260</sup>, Meghnath Dhimal<sup>261</sup>, Govinda Prasad Dhungana<sup>262</sup>, Mostafa Dianati Nasab<sup>263</sup>, Daniel Diaz<sup>264,265</sup>, Zahra Sadat Dibaji Forooshani<sup>266</sup>, Girmaye Deye Dinsa<sup>152,267</sup>, Isaac Oluwafemi Dipeolu<sup>36</sup>, Shirin Djalalinia<sup>268</sup>, Hoa Thi Do<sup>269</sup>, Huyen Phuc Do<sup>270</sup>, Paul Narh Doku<sup>271</sup>, Fariba Dorostkar<sup>272</sup>, Leila Doshmangir<sup>273</sup>, Manisha Dubey<sup>274</sup>, Bereket Duko Adema<sup>275,276</sup>, Susanna J. Dunachie<sup>277,278</sup>, Bruce B. Duncan<sup>279</sup>, Ewerton Cousin<sup>279</sup>, Andre R. Durães<sup>280,281</sup>, Lucas Earl<sup>1</sup>, Hamed Ebrahimzadeh Leylabadlo<sup>282</sup>, Aziz Eftekhari<sup>283,284</sup>, Iman El Sayed<sup>285</sup>, Maysaa El Sayed Zaki<sup>286</sup>,



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Sreeramareddy<sup>872</sup>, Rajni Kant Kant Srivastava<sup>873</sup>, Antonina Starodubova<sup>874,875</sup>, Agus Sudaryanto<sup>876,877</sup>, Mu'awiyah Babale Sufiyan<sup>878</sup>, Hafiz Ansar Rasul Suleria<sup>879</sup>, Gerhard Sulo<sup>880</sup>, Bruno F. Sunguya<sup>881,882</sup>, Bryan L. Sykes<sup>883</sup>, Rafael Tabarés-Seisdedos<sup>884,885</sup>, Takahiro Tabuchi<sup>886</sup>, Birkneh Tilahun Tadesse<sup>887,888</sup>, Amir Taherkhani<sup>889</sup>, Koku Sisay Tamirat<sup>53</sup>, Segen Gebremeskel Tassew<sup>890</sup>, Nuno Taveira<sup>891,892</sup>, Berhane Fseha Teklehaimanot<sup>893</sup>, Gebretsadkan Hintsa Tekulu<sup>894</sup>, Mohamad-Hani Tamsah<sup>895,896</sup>, Abdullah Sulieman Terkawi<sup>897,898</sup>, Zemenu Tadesse Tessema<sup>53</sup>, Nihal Thomas<sup>899</sup>, Mariya Vladimirovna Titova<sup>331,900</sup>, Kenean Getaneh Tlaye<sup>247</sup>, Hamid Reza Tohidinik<sup>420,901</sup>, Marcello Tonelli<sup>902</sup>, Marcos Roberto Tovani-Palone<sup>903</sup>, Eugenio Traini<sup>904</sup>, Khanh Bao Tran<sup>905,906</sup>, Manjari Tripathi<sup>907</sup>, Riaz Uddin<sup>496,908</sup>, Irfan Ullah<sup>909,910</sup>, Bhaskaran Unnikrishnan<sup>468</sup>, Era Upadhyay<sup>911</sup>, Ushotanefe Useh<sup>912</sup>, Muhammad Shariq Usman<sup>502</sup>, Olalekan A. Uthman<sup>913</sup>, Marco Vacante<sup>193</sup>, Masoud Vaezghasemi<sup>914</sup>, Pascual R. Valdez<sup>915,916</sup>, John VanderHeide<sup>1</sup>, Elena Varavikova<sup>917</sup>, Santosh Varughese<sup>918</sup>, Tommi Juhani Vasankari<sup>919</sup>, Yasser Vasseghian<sup>75</sup>, Yousef Veisani<sup>920</sup>, Srinivasaraghavan Venkatesh<sup>921</sup>, Narayanaswamy Venketasubramanian<sup>922,923</sup>, Madhur Verma<sup>924</sup>, Simone Vidale<sup>925</sup>, Francesco S. Violante<sup>312,926</sup>, Vasily Vlassov<sup>927</sup>, Sebastian Vollmer<sup>751,928</sup>, Rade Vukovic<sup>929,930</sup>, Yasir Waheed<sup>931</sup>, Haidong Wang<sup>1,2</sup>, Yafeng Wang<sup>932</sup>, Yuan-Pang Wang<sup>214</sup>, Girmay Teklay Weldesamuel<sup>120</sup>, Andrea Werdecker<sup>651,933</sup>, Taweewat Wiangkham<sup>934</sup>, Kirsten E. Wiens<sup>1</sup>, Tissa Wijeratne<sup>935,936</sup>, Haileab Fekadu Wolde<sup>53</sup>, Dawit Zewdu Wondafrash<sup>937,938</sup>, Tewodros Eshete Wonde<sup>490</sup>, Adam Belay Wondmienneh<sup>340,939</sup>, Ai-Min Wu<sup>940</sup>, Gelin Xu<sup>941</sup>, Abbas Yadegar<sup>614</sup>, Ali Yadollahpour<sup>942</sup>, Seyed Hossein Yahyazadeh Jabbari<sup>943</sup>, Tomohide Yamada<sup>944</sup>, Yuichiro Yano<sup>945</sup>, Sanni Yaya<sup>946</sup>, Vahid Yazdi-Feyzabadi<sup>947,948</sup>, Alex Yeshaneh<sup>949</sup>, Yigizie Yeshaw<sup>53</sup>, Yordanos Gizachew Yeshitila<sup>344</sup>, Mekdes Tigistu Yilma<sup>782</sup>, Paul Yip<sup>950,951</sup>, Naohiro Yonemoto<sup>952</sup>, Seok-Jun Yoon<sup>953</sup>, Yoosik Youm<sup>954</sup>, Mustafa Z. Younis<sup>955,956</sup>, Zabihollah Yousefi<sup>957,958</sup>, Hebat-Allah Salah A. Yousof<sup>506</sup>, Chuanhua Yu<sup>932,959</sup>, Hasan Yusefzadeh<sup>68</sup>, Telma Zahirian Moghadam<sup>69,960</sup>, Leila Zaki<sup>346</sup>, Sojib Bin Zaman<sup>431,961</sup>, Mohammad Zamani<sup>962</sup>,



**Maryam Zamanian<sup>76</sup>, Hamed Zandian<sup>960,963</sup>, Hadi Zarafshan<sup>964</sup>, Nejimu Biza Zepro<sup>184,339</sup>, Taddese Alemu Zerfu<sup>965,966</sup>, Taye Abuhay Zewale<sup>98</sup>, Yunquan Zhang<sup>967,968</sup>, Zhi-Jiang Zhang<sup>969</sup>, Xiu-Ju Zhao<sup>970</sup>, Sanjay Zodpey<sup>127</sup>, Kamiar Zomorodian<sup>706</sup>, Francis Bruno Zotor<sup>530</sup>, Ashkan Afshin<sup>1,2</sup> and Simon I. Hay<sup>1,2</sup>✉**

<sup>1</sup>Institute for Health Metrics and Evaluation, University of Washington, Seattle, WA, USA. <sup>2</sup>Department of Health Metrics Sciences, School of Medicine, University of Washington, Seattle, WA, USA. <sup>3</sup>Department of Global Health, University of Washington, Seattle, WA, USA. <sup>4</sup>Department of Medicine, University of Washington, Seattle, WA, USA. <sup>5</sup>School of Nutrition and Food Sciences, Tabriz University of Medical Sciences, Tabriz, Iran. <sup>6</sup>Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>7</sup>Advanced Diagnostic and Interventional Radiology Research Center, Tehran University of Medical Sciences, Tehran, Iran. <sup>8</sup>Department of Neurology, Cairo University, Cairo, Egypt. <sup>9</sup>Department of Parasitology and Mycology, Jahrom University of Medical Sciences, Jahrom, Iran. <sup>10</sup>The Institute of Pharmaceutical Sciences (TIPS), Toxicology and Diseases Group, Tehran University of Medical Sciences, Tehran, Iran. <sup>11</sup>Neuroscience Research Center, Isfahan University of Medical Sciences, Isfahan, Iran. <sup>12</sup>Department of Public Health, Ministry of Health, Riyadh, Saudi Arabia. <sup>13</sup>School of Public Health, Addis Ababa University, Addis Ababa, Ethiopia. <sup>14</sup>Public Health, Wachemo University, Hosanna, Ethiopia. <sup>15</sup>College of Medicine and Health Sciences, Hawassa University, Hawassa, Ethiopia. <sup>16</sup>Biostatistics and Health Informatics, Madda Walabu University, Bale Robe, Ethiopia. <sup>17</sup>Radiotherapy Center, Addis Ababa University, Addis Ababa, Ethiopia. <sup>18</sup>LABMED, Karolinska University Hospital, Huddinge, Sweden. <sup>19</sup>Research Center for Immunodeficiencies, Tehran University of Medical Sciences, Tehran, Iran. <sup>20</sup>Department of Pediatric Dentistry, Federal University of Minas Gerais, Belo Horizonte, Brazil. <sup>21</sup>Research Department, Philippine Institute for Development Studies, Quezon City, Philippines. <sup>22</sup>Cardiovascular Medicine, Ain Shams University, Abbasia, Egypt. <sup>23</sup>Bénin Clinical Research Institute (IRCB), Cotonou, Benin. <sup>24</sup>Department of Preventive Medicine, Dongguk University, Gyeongju, South Korea. <sup>25</sup>Department of Community Medicine, Kathmandu University, Devdaha, Nepal. <sup>26</sup>Hamadan University of Medical Sciences, Hamadan, Iran. <sup>27</sup>Department of Community Medicine, University of Ibadan, Ibadan, Nigeria. <sup>28</sup>Department of Community Medicine, University College Hospital, Ibadan, Ibadan, Nigeria. <sup>29</sup>Department of Sociology, Olabisi Onabanjo University, Ago-Iwoye, Nigeria. <sup>30</sup>School of Medicine, Cardiff University, Cardiff, UK. <sup>31</sup>College of Medicine, University of Ibadan, Ibadan, Nigeria. <sup>32</sup>Community Cardiovascular Research Unit, Elyon Heart Rehabilitation Center, Ibadan, Nigeria. <sup>33</sup>Department of Global Health, Stellenbosch University, Stellenbosch, South Africa. <sup>34</sup>Cochrane South Africa, South African Medical Research Council, Cape Town, South Africa. <sup>35</sup>School of Health, Ardabil University of Medical Science, Ardabil, Iran. <sup>36</sup>Department of Health Promotion and Education, University of Ibadan, Ibadan, Nigeria. <sup>37</sup>Social Behavioral Research Branch, National Institute of Health, Bethesda, MD, USA. <sup>38</sup>Cancer Prevention and Control, Georgetown University, Washington, DC, USA. <sup>39</sup>Endocrinology and Metabolism Research Center (EMRC), Tehran University of Medical Sciences, Tehran, Iran. <sup>40</sup>Epidemiology, Qom University of Medical Sciences, Qom, Iran. <sup>41</sup>Research Area for Informatics and Big Data, CSIR Institute of Genomics and Integrative Biology, Delhi, India. <sup>42</sup>Department of Internal Medicine, Baylor College of Medicine, Houston, TX, USA. <sup>43</sup>Department of Epidemiology and Health Statistics, School of Public Health, Southeast University Nanjing, Nanjing, China. <sup>44</sup>Microbiology Department, Hazara University Mansehra, Mansehra, Pakistan. <sup>45</sup>Lincoln Medical School, Universities of Nottingham & Lincoln, Lincoln, UK. <sup>46</sup>School of Advanced Technologies in Medicine, Shahid Beheshti University of Medical Sciences, Tehran, Iran. <sup>47</sup>Department of Epidemiology, Jimma University, Jimma, Ethiopia. <sup>48</sup>James P Grant School of Public Health, BRAC University, Dhaka, Bangladesh. <sup>49</sup>Health Systems and Population Studies Division, International Centre for Diarrhoeal Disease Research, Bangladesh, Dhaka, Bangladesh. <sup>50</sup>School of Community Health Sciences, University of Nevada, Reno, NV, USA. <sup>51</sup>National Malaria Elimination Program, Federal Ministry of Health, Abuja, Nigeria. <sup>52</sup>Department of Medical Laboratory Sciences, Bahir Dar University, Bahir Dar, Ethiopia. <sup>53</sup>Department of Epidemiology and Biostatistics, University of Gondar, Gondar, Ethiopia. <sup>54</sup>Department of Population Health Sciences, Duke University, Durham, NC, USA. <sup>55</sup>Duke Global Health Institute, Duke University, Durham, NC, USA. <sup>56</sup>School of Public Health and Community Medicine, University of New South Wales, Sydney, New South Wales, Australia. <sup>57</sup>John T. Milliken Department of Internal Medicine, Washington University in St. Louis, St Louis, MO, USA. <sup>58</sup>Clinical Epidemiology Center, VA Saint Louis Health Care System, Department of Veterans Affairs, St Louis, MO, USA. <sup>59</sup>Department of Medicine, Dalhousie University, Halifax, NS, Canada. <sup>60</sup>School of Health Sciences, Madda Walabu University, Bale Goba, Ethiopia. <sup>61</sup>Department of Health Information Management and Technology, Imam Abdulrahman Bin Faisal University, Dammam, Saudi Arabia. <sup>62</sup>Centre of Health System Research, National Institute of Public Health, Cuernavaca, Mexico. <sup>63</sup>Department of Pharmacy, Adigrat University, Adigrat, Ethiopia. <sup>64</sup>Medical Technical Institute, Erbil Polytechnic University, Erbil, Iraq. <sup>65</sup>Ishik University, Erbil, Iraq. <sup>66</sup>Department of Biotechnology, Quaid-i-Azam University Islamabad, Islamabad, Pakistan. <sup>67</sup>Social Determinants of Health Research Center, Qazvin University of Medical Sciences, Qazvin, Iran. <sup>68</sup>Department of Health Care Management and Economics, Urmia University of Medical Science, Urmia, Iran. <sup>69</sup>Health Management and Economics Research Center, Iran University of Medical Sciences, Tehran, Iran. <sup>70</sup>Health Economics Department, Iran University of Medical Sciences, Tehran, Iran. <sup>71</sup>Department of Microbiology, Kerman University of Medical Sciences, Kerman, Iran. <sup>72</sup>Department of Microbiology, Hormozgan University of Medical Sciences, Bandar Abbas, Iran. <sup>73</sup>Department of Health Policy and Management, Kuwait University, Safat, Kuwait. <sup>74</sup>International Centre for Casemix and Clinical Coding, National University of Malaysia, Bandar Tun Razak, Malaysia. <sup>75</sup>Research Center for Environmental Determinants of Health (RCEDH), Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>76</sup>Department of Epidemiology, Arak University of Medical Sciences, Arak, Iran. <sup>77</sup>Medical Research Center, Jazan University, Jazan, Saudi Arabia. <sup>78</sup>Department of Medical Parasitology, Sana'a University, Sana'a, Yemen. <sup>79</sup>Department of Family and Community Medicine, King Abdulaziz University, Jeddah, Saudi Arabia. <sup>80</sup>King Saud University, Riyadh, Saudi Arabia. <sup>81</sup>Research Group in Health Economics, University of Cartagena, Cartagena, Colombia. <sup>82</sup>Research Group in Hospital Management and Health Policies, University of the Coast, Barranquilla, Colombia. <sup>83</sup>Departamento de Ciencias Económicas, Universidad de la Costa, Barranquilla, Colombia. <sup>84</sup>Observatorio Nacional de Salud, National Institute of Health, Bogotá, Colombia. <sup>85</sup>Sansom Institute, South Australian Health and Medical Research Institute, Adelaide, South Australia, Australia. <sup>86</sup>Bahir Dar University, Bahir Dar, Ethiopia. <sup>87</sup>Biomedical Science, University of Cape Coast, Cape Coast, Ghana. <sup>88</sup>Health Services Management Department, Arak University of Medical Sciences, Arak, Iran. <sup>89</sup>Health Services Management, Isfahan University of Medical Sciences, Isfahan, Iran. <sup>90</sup>Department of Radiology, Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>91</sup>Department of Epidemiology and Biostatistics, University of the Philippines Manila, Manila, Philippines. <sup>92</sup>Online Programs for Applied Learning, Johns Hopkins University, Baltimore, MD, USA. <sup>93</sup>Mansoura University, Mansoura, Egypt. <sup>94</sup>Carol Davila University of Medicine and Pharmacy, Bucharest, Romania. <sup>95</sup>Research Center for Evidence Based Medicine-Health Management and Safety Promotion Research Institute, Tabriz University of Medical Sciences, Tabriz, Iran. <sup>96</sup>Razi Vaccine and Serum Research Institute, Agricultural Research Education and Extension Organization (AREEO), Tehran, Iran. <sup>97</sup>Department of Epidemiology and Biostatistics, Health Promotion Research Center, Zahedan, Iran. <sup>98</sup>Department of Epidemiology and Biostatistics, Bahir Dar University, Bahir Dar, Ethiopia. <sup>99</sup>Department of Health Policy and Administration, University of the Philippines Manila, Manila, Philippines. <sup>100</sup>Department of Applied Social Sciences, Hong Kong Polytechnic University, Hong Kong, China. <sup>101</sup>Department of Agribusiness, Universitas Sebelas Maret, Surakarta, Indonesia. <sup>102</sup>Department of Parasitology, Mazandaran University of Medical Sciences, Sari, Iran. <sup>103</sup>Department of Microbiology and Immunology, Iranshahr University of Medical Sciences, Iranshahr, Iran. <sup>104</sup>Department of Pathology, Al-Imam Mohammad Ibn Saud Islamic University, Riyadh, Saudi Arabia. <sup>105</sup>Department of Sociology and Social Work, Kwame Nkrumah

University of Science and Technology, Kumasi, Ghana. <sup>106</sup>Center for International Health, Ludwig Maximilians University, Munich, Germany. <sup>107</sup>Social Determinants of Health Research Center, Birjand University of Medical Sciences, Birjand, Iran. <sup>108</sup>Department of Health Promotion and Education, Tehran University of Medical Sciences, Tehran, Iran. <sup>109</sup>School of Health Sciences, Birmingham City University, Birmingham, UK. <sup>110</sup>Department of Neurobiology, Karolinska Institutet, Stockholm, Sweden. <sup>111</sup>School of Health and Social Studies, Dalarna University, Falun, Sweden. <sup>112</sup>School of Nursing and Midwife, Babol University of Medical Sciences, Babol, Iran. <sup>113</sup>Babol University of Medical Sciences, Babol, Iran. <sup>114</sup>Preventive Medicine and Public Health Research Center, Iran University of Medical Sciences, Tehran, Iran. <sup>115</sup>Neurology, Shiraz University of Medical Sciences, Shiraz, Iran. <sup>116</sup>Prevention of Metabolic Disorders Research Center, Shahid Beheshti University of Medical Sciences, Tehran, Iran. <sup>117</sup>Department of Microbiology, Hamedan University of Medical Sciences, Hamedan, Iran. <sup>118</sup>Department of Clinical Chemistry, University of Gondar, Gondar, Ethiopia. <sup>119</sup>Non-Communicable Diseases Research Center, Tehran University of Medical Sciences, Tehran, Iran. <sup>120</sup>Department of Nursing, Aksum University, Aksum, Ethiopia. <sup>121</sup>Department of Health System and Health Economics, Bahir Dar University, Bahir Dar City, Ethiopia. <sup>122</sup>School of Nursing, , University of Nottingham, Amman, Jordan. <sup>123</sup>School of Business, University of Leicester, Leicester, UK. <sup>124</sup>Department of Statistics and Econometrics, Bucharest University of Economic Studies, Bucharest, Romania. <sup>125</sup>Bénin Clinical Research Institute (IRCB), Abomey-Calavi, Benin. <sup>126</sup>Contrôle des Maladies Infectieuses, Laboratory of Studies and Research-Action in Health, Porto Novo, Benin. <sup>127</sup>Indian Institute of Public Health, Public Health Foundation of India, Gurugram, India. <sup>128</sup>The Judith Lumley Centre, La Trobe University, Melbourne, Victoria, Australia. <sup>129</sup>General Office for Research and Technological Transfer, Peruvian National Institute of Health, Lima, Peru. <sup>130</sup>Department of Health Policy Planning and Management, University of Health and Allied Sciences, Ho, Ghana. <sup>131</sup>Department of Nursing, Debre Berhan University, Debre Berhan, Ethiopia. <sup>132</sup>Cellular and Molecular Biology Research Center, Babol University of Medical Sciences, Babol, Iran. <sup>133</sup>Department of Environmental Health Engineering, Hamadan University of Medical Sciences, Hamadan, Iran. <sup>134</sup>Department of Reproductive Health, University of Gondar, Gondar, Ethiopia. <sup>135</sup>Public Health Risk Sciences Division, Public Health Agency of Canada, Toronto, Ontario, Canada. <sup>136</sup>Department of Nutritional Sciences, University of Toronto, Toronto, Ontario, Canada. <sup>137</sup>Department of Forensic Science, Government Institute of Forensic Science, Nagpur, India. <sup>138</sup>Healthcare Management Department, Shiraz University of Medical Sciences, Shiraz, Iran. <sup>139</sup>Biochemistry Unit, Universiti Sultan Zainal Abidin, Kuala Terengganu, Malaysia. <sup>140</sup>Biomedicine Department, Universiti Sultan Zainal Abidin Gongbedak, Kuala Terengganu, Malaysia. <sup>141</sup>Health Policy and Management Department, Tehran University of Medical Sciences, Tehran, Iran. <sup>142</sup>Department of Forensic Medicine and Toxicology, , Manipal Academy of Higher Education, Manipal, India. <sup>143</sup>Department of Medical Laboratory Science, Haramaya University, Harar, Ethiopia. <sup>144</sup>School of Public Health, Haramaya University, Harar, Ethiopia. <sup>145</sup>Department of Hypertension, Medical University of Lodz, Lodz, Poland. <sup>146</sup>Polish Mothers' Memorial Hospital Research Institute, Lodz, Poland. <sup>147</sup>Department of Noncommunicable Diseases, Bangladesh University of Health Sciences (BUHS), Dhaka, Bangladesh. <sup>148</sup>Department of Animal Pathology and Epidemiology, Razi Vaccine and Serum Research Institute, Karaj, Iran. <sup>149</sup>Department of Neurosciences, Costa Rican Department of Social Security, San Jose, Costa Rica. <sup>150</sup>School of Medicine, University of Costa Rica, San Pedro, Costa Rica. <sup>151</sup>Heidelberg Institute of Global Health (HIGH), Heidelberg University, Heidelberg, Germany. <sup>152</sup>T.H. Chan School of Public Health, Harvard University, Boston, MA, USA. <sup>153</sup>University of Aden, Aden, Yemen. <sup>154</sup>School of Public Health, Imperial College London, London, UK. <sup>155</sup>Health Human Resources Research Center, Shiraz University of Medical Sciences, Shiraz, Iran. <sup>156</sup>Department of Applied Human Nutrition, Bahir Dar University, Bahir Dar, Ethiopia. <sup>157</sup>University of Gondar, Gondar, Ethiopia. <sup>158</sup>Department of Community Medicine, Gandhi Medical College Bhopal, Bhopal, India. <sup>159</sup>Jazan University, Jazan, Saudi Arabia. <sup>160</sup>Social Determinants of Health Research Center, Lorestan University of Medical Sciences, Khorramabad, Iran. <sup>161</sup>Department of Epidemiology and Biostatistics, Lorestan University of Medical Sciences, Khorramabad, Iran. <sup>162</sup>Department of Reproductive Health and Population Studies, Bahir Dar University, Bahir Dar, Ethiopia. <sup>163</sup>Nuffield Department of Population Health, University of Oxford, Oxford, UK. <sup>164</sup>Department of Public Health, Arba Minch University, Arba Minch, Ethiopia. <sup>165</sup>Department of Nutrition and Dietetics, Mekelle University, Mekelle, Ethiopia. <sup>166</sup>Adigrat University, Adigrat, Ethiopia. <sup>167</sup>School of Public Health, Wolaita Sodo University, Addis Ababa, Ethiopia. <sup>168</sup>Department of Medicine, Medical College of Georgia at Augusta University, Augusta, GA, USA. <sup>169</sup>Hubert Department of Global Health, Emory University, Atlanta, GA, USA. <sup>170</sup>Department of Global Health, University of South Florida, Tampa, FL, USA. <sup>171</sup>Department of Health Information Management, Manipal Academy of Higher Education, Manipal, Manipal, India. <sup>172</sup>School of Public Health, University of Adelaide, Adelaide, South Australia, Australia. <sup>173</sup>Public Health Research Laboratory, Institute of Medicine, Tribhuvan University, Kathmandu, Nepal. <sup>174</sup>Department of Community Medicine and Family Medicine, All India Institute of Medical Sciences, Jodhpur, India. <sup>175</sup>Department of Community Medicine, Datta Meghe Institute of Medical Sciences, Deemed University, Wardha, India. <sup>176</sup>Department of Statistical and Computational Genomics, National Institute of Biomedical Genomics, Kalyani, India. <sup>177</sup>Department of Statistics, University of Calcutta, Kolkata, India. <sup>178</sup>Social Determinants of Health Research Center, Babol University of Medical Sciences, Babol, Iran. <sup>179</sup>Istituto di Ricerche Farmacologiche Mario Negri IRCCS, Ranica, Italy. <sup>180</sup>Health Economics & Outcomes Research, Creativ-Ceutical (Huntsworth Health), London, UK. <sup>181</sup>Woldia University, Woldia, Ethiopia. <sup>182</sup>National Centre for Epidemiology and Population Health, Australian National University, Canberra, Australian Capital Territory, Australia. <sup>183</sup>Department of Clinical Pharmacy and Pharmacology, University of Dhaka, Dhaka, Bangladesh. <sup>184</sup>Department of Public Health, Samara University, Samara, Ethiopia. <sup>185</sup>Debre Tabor University, Addis Ababa University, Debre Tabor, Ethiopia. <sup>186</sup>Department of Pediatrics and Child Health Nursing, Bahir Dar University, Bahir Dar, Ethiopia. <sup>187</sup>Transport and Road Safety (TARS) Research Center, , University of New South Wales, Sydney, New South Wales, Australia. <sup>188</sup>School of Health Sciences, Swinburne University of Technology, Melbourne, Victoria, Australia. <sup>189</sup>Department of Nutrition, Saint Paul's Hospital Millennium Medical College, Addis Ababa, Ethiopia. <sup>190</sup>Department of Veterinary Medicine, Islamic Azad University, Kermanshah, Iran. <sup>191</sup>Department of Biomedical Sciences, Nazarbayev University, Nur-Sultan City, Kazakhstan. <sup>192</sup>Department of Internal Medicine, Manipal Academy of Higher Education, Mangalore, India. <sup>193</sup>Department of General Surgery and Medical-Surgical Specialties, University of Catania, Catania, Italy. <sup>194</sup>School of Medicine, Hamadan University of Medical Sciences, Hamadan, Iran. <sup>195</sup>Department of Infectious Disease Epidemiology, London School of Hygiene & Tropical Medicine, London, UK. <sup>196</sup>University of Genoa, Genoa, Italy. <sup>197</sup>Division of Hematology and Oncology, Georgetown University, Washington, DC, USA. <sup>198</sup>Epidemiology and Evidence Based Medicine, I.M. Sechenov First Moscow State Medical University, Moscow, Russia. <sup>199</sup>Gorgas Memorial Institute for Health Studies, Panama City, Panama. <sup>200</sup>Department of Research, Golden Community, Kathmandu, Nepal. <sup>201</sup>Department of Community Medicine, Employees' State Insurance Model Hospital, Bangalore, India. <sup>202</sup>Department for Health Care Management, Technical University of Berlin, Berlin, Germany. <sup>203</sup>School of Public Health and Health Systems, University of Waterloo, Waterloo, Ontario, Canada. <sup>204</sup>Al Shifa School of Public Health, Al Shifa Trust Eye Hospital, Rawalpindi, Pakistan. <sup>205</sup>Internal Medicine Department, Hospital Italiano de Buenos Aires, Ciudad Autónoma de Buenos Aires, Buenos Aires, Argentina. <sup>206</sup>Comisión Directiva, Argentine Society of Medicine, Ciudad Autónoma de Buenos Aires, Buenos Aires, Argentina. <sup>207</sup>National Institute of Public Health, Cuernavaca, Mexico. <sup>208</sup>Department of Disease Control, London School of Hygiene & Tropical Medicine, London, UK. <sup>209</sup>Centre for Population Health Sciences, Nanyang Technological University, Singapore, Singapore. <sup>210</sup>Global eHealth Unit, Imperial College London, London, UK. <sup>211</sup>Department of Population and Health, Metropolitan Autonomous University, Mexico City, Mexico. <sup>212</sup>Department of Medical Epidemiology and Biostatistics, Karolinska Institutet, Stockholm, Sweden. <sup>213</sup>Research Unit on Applied Molecular Biosciences (UCIBIO), University of Porto, Porto, Portugal. <sup>214</sup>Department of Psychiatry, University of São Paulo, São Paulo, Brazil. <sup>215</sup>Colombian National Health Observatory, National Institute of Health, Bogota, Colombia. <sup>216</sup>Epidemiology and Public Health Evaluation Group, National University of Colombia, Bogota, Colombia. <sup>217</sup>Mary MacKillop Institute for Health Research, Australian Catholic University, Melbourne, Victoria, Australia. <sup>218</sup>School of Public Health, University of Hong Kong, Hong Kong, China. <sup>219</sup>Health, Nutrition and Population, World Bank, Lusaka, Zambia. <sup>220</sup>Institute for Global Health, Heidelberg University, Heidelberg, Germany. <sup>221</sup>Department of Pharmacology, All India Institute of Medical Sciences, Jodhpur, India. <sup>222</sup>Division of Epidemiology, National Institute of Cholera and Enteric Diseases, Kolkata, India. <sup>223</sup>Department of Medicine, University of Toronto, Toronto,

Ontario, Canada. <sup>224</sup>Population Research Centre, Gokhale Institute of Politics and Economics, Pune, India. <sup>225</sup>International Institute for Population Sciences, Mumbai, India. <sup>226</sup>Department of Medical Entomology and Vector Control, Urmia University of Medical Science, Urmia, Iran. <sup>227</sup>Department of Biostatistics and Epidemiology, Babol University of Medical Sciences, Babol, Iran. <sup>228</sup>Epidemiology Research Center, Royan Institute, Tehran, Iran. <sup>229</sup>Department of Nursing, Wolaita Sodo University, Sodo, Ethiopia. <sup>230</sup>Department of Epidemiology and Preventive Medicine, Monash University, Melbourne, Victoria, Australia. <sup>231</sup>Department of Pulmonary Medicine, Christian Medical College and Hospital (CMC), Vellore, India. <sup>232</sup>Hanoi National University of Education, Hanoi, Vietnam. <sup>233</sup>School of Public Health and Preventive Medicine, Monash University, Melbourne, Victoria, Australia. <sup>234</sup>School of Medicine and Surgery, University of Milan Bicocca, Monza, Italy. <sup>235</sup>Institute of Public Health, University of Gondar, Gondar, Ethiopia. <sup>236</sup>Discipline of Public Health, Flinders University, Adelaide, South Australia, Australia. <sup>237</sup>Department of Human Physiology, University of Gondar, Gondar, Ethiopia. <sup>238</sup>Department of Environmental Health, Imam Abdulrahman Bin Faisal University, Dammam, Saudi Arabia. <sup>239</sup>Department of Dermatology, Case Western Reserve University, Cleveland, OH, USA. <sup>240</sup>Department of Dermatology, University of Milan, Milan, Italy. <sup>241</sup>Department of Pediatrics, Tanta University, Tanta, Egypt. <sup>242</sup>Toxoplasmosis Research Center, Mazandaran University of Medical Sciences, Sari, Iran. <sup>243</sup>Division of Women and Child Health, Aga Khan University, Karachi, Pakistan. <sup>244</sup>Department of Epidemiology and Biostatistics, Arnold School of Public Health, University of South Carolina, Columbia, SC, USA. <sup>245</sup>Population and Development, Facultad Latinoamericana de Ciencias Sociales Mexico, Mexico City, Mexico. <sup>246</sup>Australian Institute for Suicide Research and Prevention, Griffith University, Mount Gravatt, Queensland, Australia. <sup>247</sup>Department of Nursing, Woldia University, Woldia, Ethiopia. <sup>248</sup>Department of Nursing, Jimma University, Jimma, Ethiopia. <sup>249</sup>Department of Neonatal Nursing, St. Paul's Hospital Millennium Medical College, Addis Ababa, Ethiopia. <sup>250</sup>Ambo University, Ambo, Ethiopia. <sup>251</sup>School of Pharmacy, Aksum University, Aksum, Ethiopia. <sup>252</sup>Addis Ababa University, Addis Ababa, Ethiopia. <sup>253</sup>Center for Nutrition and Health Research, National Institute of Public Health, Cuernavaca, Mexico. <sup>254</sup>Department of Global Health and Infection, Brighton and Sussex Medical School, Brighton, UK. <sup>255</sup>Division of Cardiology, Atlanta Veterans Affairs Medical Center, Decatur, GA, USA. <sup>256</sup>School of Nutrition, Food Science and Technology, Hawassa University, Hawassa, Ethiopia. <sup>257</sup>School of Nursing and Midwifery, Haramaya University, Harar, Ethiopia. <sup>258</sup>Centre for Atmospheric Sciences, Indian Institute of Technology Delhi, New Delhi, India. <sup>259</sup>Department of Community Medicine, University of Peradeniya, Peradeniya, Sri Lanka. <sup>260</sup>Mathematical Demography and Statistics, International Institute for Population Sciences, Mumbai, India. <sup>261</sup>Health Research Section, Nepal Health Research Council, Kathmandu, Nepal. <sup>262</sup>Department of Microbiology, Far Western University, Mahendranagar, Nepal. <sup>263</sup>Department of Epidemiology, Shiraz University of Medical Sciences, Shiraz, Iran. <sup>264</sup>Center of Complexity Sciences, National Autonomous University of Mexico, Mexico City, Mexico. <sup>265</sup>Facultad de Medicina Veterinaria y Zootecnia, Autonomous University of Sinaloa, Culiacan, Mexico. <sup>266</sup>Department of Nursing, Bank Mellé, Tehran, Iran. <sup>267</sup>Fenot Project, Harvard University, Addis Ababa, Ethiopia. <sup>268</sup>Ministry of Health and Medical Education, Tehran, Iran. <sup>269</sup>Center of Excellence in Public Health Nutrition, Nguyen Tat Thanh University, Ho Chi Minh City, Vietnam. <sup>270</sup>Center of Excellence in Behavioral Medicine, Nguyen Tat Thanh University, Ho Chi Minh City, Vietnam. <sup>271</sup>School of Nursing and Midwifery, University of Cape Coast, Cape Coast, Ghana. <sup>272</sup>Iran University of Medical Sciences, Tehran, Iran. <sup>273</sup>Department of Health Policy and Economy, Tabriz University of Medical Sciences, Tabriz, Iran. <sup>274</sup>World Food Programme, New Delhi, India. <sup>275</sup>Public Health Department, Hawassa University, Hawassa, Ethiopia. <sup>276</sup>Curtin University, Perth, Western Australia, Australia. <sup>277</sup>Centre for Tropical Medicine and Global Health, University of Oxford, Oxford, UK. <sup>278</sup>Mahidol-Oxford Tropical Medicine Research Unit, Bangkok, Thailand. <sup>279</sup>Postgraduate Program in Epidemiology, Federal University of Rio Grande do Sul, Porto Alegre, Brazil. <sup>280</sup>School of Medicine, Federal University of Bahia, Salvador, Brazil. <sup>281</sup>Medicina Interna, Escola Bahiana de Medicina e Saúde Pública, Salvador, Brazil. <sup>282</sup>Department of Bacteriology and Virology, Tabriz University of Medical Sciences, Tabriz, Iran. <sup>283</sup>Department of Pharmacology and Toxicology, Maragheh University of Medical Sciences, Maragheh, Iran. <sup>284</sup>Department of Pharmacology and Toxicology, Tabriz University of Medical Sciences, Tabriz, Iran. <sup>285</sup>Biomedical Informatics and Medical Statistics, Alexandria University, Alexandria, Egypt. <sup>286</sup>Department of Clinical Pathology, Mansoura University, Mansoura, Egypt. <sup>287</sup>Pediatric Dentistry and Dental Public Health, Alexandria University, Alexandria, Egypt. <sup>288</sup>Institute of Public Health, United Arab Emirates University, Al Ain, United Arab Emirates. <sup>289</sup>Department of Statistics, Debre Markos University, Debre Markos, Ethiopia. <sup>290</sup>Department of Public Health Sciences, Karolinska Institutet, Stockholm, Sweden. <sup>291</sup>World Health Programme, Université du Québec en Abitibi-Témiscamingue, Rouyn-Noranda, Quebec, Canada. <sup>292</sup>Endemic Medicine and Hepatogastroenterology Department, Cairo University, Cairo, Egypt. <sup>293</sup>Department of Biosciences, Nottingham Trent University, Nottingham, UK. <sup>294</sup>Eijkman-Oxford Clinical Research Unit, Eijkman Institute for Molecular Biology, Jakarta, Indonesia. <sup>295</sup>Ophthalmic Epidemiology Research Center, Shahroud University of Medical Sciences, Shahroud, Iran. <sup>296</sup>Department of Microbiology and Immunology, Suez Canal University, Ismailia, Egypt. <sup>297</sup>Department of Midwifery, Wolkite University, Wolkite, Ethiopia. <sup>298</sup>Department of Midwifery, Woldia University, Woldia, Ethiopia. <sup>299</sup>Department of Medicinal Chemistry, Kerman University of Medical Sciences, Kerman, Iran. <sup>300</sup>Pharmaceutics Research Center, Kerman University of Medical Sciences, Kerman, Iran. <sup>301</sup>Multiple Sclerosis Research Center, Tehran University of Medical Sciences, Tehran, Iran. <sup>302</sup>Department of Physiology, Tarbiat Modares University, Tehran, Iran. <sup>303</sup>Division of Cancer Epidemiology and Genetics, National Cancer Institute, Bethesda, MD, USA. <sup>304</sup>Tehran University of Medical Sciences, Tehran, Iran. <sup>305</sup>Unit of Medical Physiology, Hawassa University, Hawassa, Ethiopia. <sup>306</sup>Berman Institute of Bioethics, Johns Hopkins University, Baltimore, MD, USA. <sup>307</sup>Nutrition and Food Systems Division, Food and Agriculture Organization of the United Nations, Rome, Italy. <sup>308</sup>School of Public Health, Tehran University of Medical Sciences, Tehran, Iran. <sup>309</sup>Department of Political Science, University of Human Development, Sulaimaniyah, Iraq. <sup>310</sup>Deputy of Research and Technology, Hamadan University of Medical Sciences, Hamadan, Iran. <sup>311</sup>College of Medicine, Imam Mohammad Ibn Saud Islamic University, Riyadh, Saudi Arabia. <sup>312</sup>Department of Medical and Surgical Sciences, University of Bologna, Bologna, Italy. <sup>313</sup>Department of Psychology, Federal University of Sergipe, Sao Cristovao, Brazil. <sup>314</sup>Department of Biological and Biomedical Sciences, Aga Khan University, Karachi, Pakistan. <sup>315</sup>College of Medicine and Public Health, Flinders University, Adelaide, South Australia, Australia. <sup>316</sup>Institute of Resource Governance and Social Change, Kupang, Indonesia. <sup>317</sup>Social Determinants of Health Research Center, Hamadan University of Medical Sciences, Hamadan, Iran. <sup>318</sup>Department of Public Health Nutrition, Bahir Dar University, Bahir Dar, Ethiopia. <sup>319</sup>School of Nursing and Midwifery, Hawassa University, Hawassa, Ethiopia. <sup>320</sup>Division of Neurology, University of Ottawa, Ottawa, Ontario, Canada. <sup>321</sup>REQUIMTE/LAQV - Network of Chemistry and Technology, University of Porto, Porto, Portugal. <sup>322</sup>Center for Biotechnology and Fine Chemistry, Catholic University of Portugal, Porto, Portugal. <sup>323</sup>Department of Health Education & Behavioral Sciences, Jimma University, Jimma, Ethiopia. <sup>324</sup>Jimma University, Jimma, Ethiopia. <sup>325</sup>Psychiatry Department, Kaiser Permanente, Fontana, CA, USA. <sup>326</sup>School of Health Sciences, A.T. Still University, Mesa, AZ, USA. <sup>327</sup>Department of Population Medicine and Health Services Research, Bielefeld University, Bielefeld, Germany. <sup>328</sup>Unit for Population-Based Dermatology Research, King's College London, London, UK. <sup>329</sup>Institute of Gerontology, National Academy of Medical Sciences of Ukraine, Kyiv, Ukraine. <sup>330</sup>Department of Child Dental Health, Obafemi Awolowo University, Ile-Ife, Nigeria. <sup>331</sup>Timiryazev Institute of Plant Physiology (IPPRAS), Russian Academy of Sciences, Moscow, Russia. <sup>332</sup>Abadan School of Medical Sciences, Abadan University of Medical Sciences, Abadan, Iran. <sup>333</sup>Department of Research, Center for Population and Health, Wiesbaden, Germany. <sup>334</sup>Department of Family Medicine and Primary Care, University of the Witwatersrand, Johannesburg, South Africa. <sup>335</sup>Department of Dermatology, Kobe University, Kobe, Japan. <sup>336</sup>Gene Expression & Regulation Program, The Wistar Institute, Philadelphia, PA, USA. <sup>337</sup>School of Nursing and Midwifery, Wollega University, Nekemte, Ethiopia. <sup>338</sup>Public Health Department, Madda Walabu University, Bale-Robe, Ethiopia. <sup>339</sup>School of Public Health, Mekelle University, Mekelle, Ethiopia. <sup>340</sup>Department of Nursing and Midwifery, Addis Ababa University, Addis Ababa, Ethiopia. <sup>341</sup>Nursing Department, Mekelle University, Mekelle, Ethiopia. <sup>342</sup>Haramaya University, Dire Dawa, Ethiopia. <sup>343</sup>Pharmacy, Wollo University, Dessie, Ethiopia. <sup>344</sup>Department of Nursing, Arba Minch University, Arba Minch, Ethiopia. <sup>345</sup>Department of Biostatistics, Mekelle University, Mekelle, Ethiopia. <sup>346</sup>Department of Parasitology and Entomology, Tarbiat Modares University, Tehran, Iran. <sup>347</sup>Department of Medical Surgery, Tabriz University of Medical Sciences, Tabriz, Iran. <sup>348</sup>Department of Medicine,



Massachusetts General Hospital, Boston, MA, USA. <sup>349</sup>Neuroscience Institute, Academy of Medical Science, Tehran, Iran. <sup>350</sup>Department of Health Services Management, Iran University of Medical Sciences, Tehran, Iran. <sup>351</sup>Social Determinants of Health Research Center, University of Social Welfare and Rehabilitation Sciences, Tehran, Iran. <sup>352</sup>Science and Research Branch, Islamic Azad University, Tehran, Iran. <sup>353</sup>Young Researchers and Elite Club, Islamic Azad University, Rasht, Iran. <sup>354</sup>University of Lahore, Lahore, Pakistan. <sup>355</sup>Afro-Asian Institute, Lahore, Pakistan. <sup>356</sup>Adelaide Medical School, University of Adelaide, Adelaide, South Australia, Australia. <sup>357</sup>Department of Family and Community Medicine, University of Hail, Hail, Saudi Arabia. <sup>358</sup>Center for the Study of Regional Development, Jawahar Lal Nehru University, New Delhi, India. <sup>359</sup>Department of Chemistry, University of Porto, Porto, Portugal. <sup>360</sup>Department of Biostatistics and Epidemiology, University of Oklahoma, Oklahoma City, OK, USA. <sup>361</sup>Department of Health and Social Affairs, Government of the Federated States of Micronesia, Palikir, Federated States of Micronesia. <sup>362</sup>Department of Respiratory Medicine, Hokkaido University, Sapporo, Japan. <sup>363</sup>Center for Environmental and Health Sciences, Hokkaido University, Sapporo, Japan. <sup>364</sup>Center for Clinical and Epidemiological Research, University of São Paulo, São Paulo, Brazil. <sup>365</sup>Internal Medicine Department, University of São Paulo, São Paulo, Brazil. <sup>366</sup>Manipal Institute of Virology, Manipal Academy of Higher Education, Manipal, India. <sup>367</sup>Department of Dermatology, Boston University, Boston, MA, USA. <sup>368</sup>Instituto de Patologia Tropical e Saúde Pública, Federal University of Goiás, Goiânia, Brazil. <sup>369</sup>College of Medicine and Health Science, Jigjiga University, Jigjiga, Ethiopia. <sup>370</sup>Department of Epidemiology and Biostatistics, Zhengzhou University, Zhengzhou, China. <sup>371</sup>March of Dimes, Arlington, VA, USA. <sup>372</sup>School of Public Health, West Virginia University Morgantown, Morgantown, WV, USA. <sup>373</sup>Academics and Research Department, Rajasthan University of Health Sciences, Jaipur, India. <sup>374</sup>Department of Medicine, Mahatma Gandhi University of Medical Sciences & Technology, Jaipur, India. <sup>375</sup>Department of Radiology and Radiological Sciences, Johns Hopkins University, Baltimore, MD, USA. <sup>376</sup>School of Medicine, Tehran University of Medical Sciences, Tehran, Iran. <sup>377</sup>Department of Nursing, St. Paul's Hospital Millennium Medical College, Addis Ababa, Ethiopia. <sup>378</sup>Department of Pharmacology, Tehran University of Medical Sciences, Tehran, Iran. <sup>379</sup>Obesity Research Center, Shahid Beheshti University of Medical Sciences, Tehran, Iran. <sup>380</sup>Global and Community Mental Health Research Group, University of Macau, Macao, China. <sup>381</sup>Department of Anatomical Sciences, Tarbiat Modares University, Tehran, Iran. <sup>382</sup>Department of Family and Community Medicine, Arabian Gulf University, Manama, Bahrain. <sup>383</sup>Department of Health Management and Economics, Hamadan University of Medical Sciences, Hamadan, Iran. <sup>384</sup>School of Medicine, University of Western Australia, Perth, Western Australia, Australia. <sup>385</sup>Neurology Department, Sir Charles Gairdner Hospital, Perth, Western Australia, Australia. <sup>386</sup>Tabriz University of Medical Sciences, Tabriz, Iran. <sup>387</sup>Department of Dental Public Health, Universitas Airlangga Indonesia, Surabaya, Indonesia. <sup>388</sup>Australian Research Centre for Population Oral Health, University of Adelaide, Adelaide, South Australia, Australia. <sup>389</sup>Department of Zoology, Al-Azhar University, Cairo, Egypt. <sup>390</sup>Institute for Social Science Research, The University of Queensland, Indooroopilly, Queensland, Australia. <sup>391</sup>Department of Healthcare Management, Maragheh University of Medical Sciences, Maragheh, Iran. <sup>392</sup>Department of Microbiology, Maragheh University of Medical Sciences, Maragheh, Iran. <sup>393</sup>Department of Microbiology, Tehran University of Medical Sciences, Tehran, Iran. <sup>394</sup>Department of Biology, Utica College, Utica, NY, USA. <sup>395</sup>Gastrointestinal and Liver Disease Research Center, Guilan University of Medical Sciences, Rasht, Iran. <sup>396</sup>Guilan University of Medical Sciences, Rasht, Iran. <sup>397</sup>Department of Public Health, Mizan-Tepi University, Tepi, Ethiopia. <sup>398</sup>Unit of Epidemiology and Social Medicine, University Hospital Antwerp, Wilrijk, Belgium. <sup>399</sup>Department of Clinical Sciences, Karolinska University Hospital, Stockholm, Sweden. <sup>400</sup>School of Health Sciences, City University of London, London, UK. <sup>401</sup>Institute of Pharmaceutical Sciences, University of Veterinary and Animal Sciences, Lahore, Pakistan. <sup>402</sup>Department of Pharmacy Administration and Clinical Pharmacy, Xian Jiaotong University, Xian, China. <sup>403</sup>Shahrekord University of Medical Sciences, Shahrekord, Iran. <sup>404</sup>School of Public Health, Curtin University, Perth, Western Australia, Australia. <sup>405</sup>Agriculture and Food, Commonwealth Scientific and Industrial Research Organisation, St. Lucia, Queensland, Australia. <sup>406</sup>Medical Biology Research Center, Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>407</sup>Department of Biostatistics and Epidemiology, Adigrat University, Adigrat, Ethiopia. <sup>408</sup>Department of Psychiatry, University Medical Center Groningen, Groningen, the Netherlands. <sup>409</sup>Department of Epidemiology, Columbia University, New York, NY, USA. <sup>410</sup>Department of Pediatrics, Dell Medical School, University of Texas Austin, Austin, TX, USA. <sup>411</sup>Kasturba Medical College, Manipal Academy of Higher Education, Manipal, India. <sup>412</sup>Guilan Road Trauma Research Center, Guilan University of Medical Sciences, Rasht, Iran. <sup>413</sup>Social Determinants of Health Research Center, Guilan University of Medical Sciences, Rasht, Iran. <sup>414</sup>Department of Pediatrics, Yonsei University, Seoul, South Korea. <sup>415</sup>Research Department, Electronic Medical Records for the Developing World, York, UK. <sup>416</sup>Transdisciplinary Centre for Qualitative Methods, Manipal Academy of Higher Education, Manipal, India. <sup>417</sup>Nevada Division of Public and Behavioral Health, Carson City, NV, USA. <sup>418</sup>Department of Pharmacology and Therapeutics, Dhaka Medical College, Dhaka, Bangladesh. <sup>419</sup>Department of Pharmacology, Bangladesh Industrial Gases Limited, Tangail, Bangladesh. <sup>420</sup>Department of Epidemiology and Biostatistics, Tehran University of Medical Sciences, Tehran, Iran. <sup>421</sup>Department of Computer Engineering, Islamic Azad University, Tehran, Iran. <sup>422</sup>Computer Science Department, University of Human Development, Sulaymaniyah, Iraq. <sup>423</sup>Department of General Surgery, Carol Davila University of Medicine and Pharmacy, Bucharest, Romania. <sup>424</sup>Department of Internal Medicine, Bucharest Emergency Hospital, Bucharest, Romania. <sup>425</sup>Faculty of Dentistry, Department of Legal Medicine and Bioethics, Carol Davila University of Medicine and Pharmacy, Bucharest, Romania. <sup>426</sup>Clinical Legal Medicine Department, National Institute of Legal Medicine, Bucharest, Romania. <sup>427</sup>College of Science and Engineering, Hamad Bin Khalifa University, Doha, Qatar. <sup>428</sup>Medicine School of Tunis, Baab Saadoun, Tunisia. <sup>429</sup>Department of Epidemiology and Health Statistics, Central South University, Changsha, China. <sup>430</sup>School of Public Health, University of Sydney, Sydney, New South Wales, Australia. <sup>431</sup>Maternal and Child Health Division, International Centre for Diarrhoeal Disease Research, Dhaka, Bangladesh. <sup>432</sup>Department of Public Health and Community Medicine, Shaikh Khalifa Bin Zayed Al-Nahyan Medical College at Shaikh Zayed Medical Complex, Lahore, Pakistan. <sup>433</sup>Department of Occupational Safety and Health, China Medical University, Taichung, Taiwan. <sup>434</sup>Department of Epidemiology, University of Kragujevac, Kragujevac, Serbia. <sup>435</sup>Department of Public Health, Lorestan University of Medical Sciences, Khorramabad, Iran. <sup>436</sup>Department of Family Medicine, Bangalore Baptist Hospital, Bangalore, India. <sup>437</sup>Global Health and Development Department, Taipei Medical University, Taipei City, Taiwan. <sup>438</sup>Research Institute for Endocrine Sciences, Shahid Beheshti University of Medical Sciences, Tehran, Iran. <sup>439</sup>Institute for Physical Activity and Nutrition, Deakin University, Burwood, Victoria, Australia. <sup>440</sup>Sydney Medical School, University of Sydney, Sydney, New South Wales, Australia. <sup>441</sup>School of Health Systems and Public Health, University of Pretoria, Hatfield, South Africa. <sup>442</sup>Cochrane Center, South African Medical Research Council, Parow Valley, South Africa. <sup>443</sup>Health Systems and Public Health, Stellenbosch University, Cape Town, South Africa. <sup>444</sup>Department of Epidemiology, Shahid Beheshti University of Medical Sciences, Tehran, Iran. <sup>445</sup>Department of Environmental Health Engineering, Guilan University of Medical Sciences, Rasht, Iran. <sup>446</sup>Medical Research Council South Africa, Cape Town, South Africa. <sup>447</sup>Centre for Evidence Based Health Care, Stellenbosch University, Cape Town, South Africa. <sup>448</sup>Department of Immunology, Tabriz University of Medical Sciences, Tabriz, Iran. <sup>449</sup>Department of Psychosis, Babol Noshirvani University of Technology, Babol, Iran. <sup>450</sup>Department of Immunology, Isfahan University of Medical Sciences, Isfahan, Iran. <sup>451</sup>Department for Health Care and Public Health, Sechenov First Moscow State Medical University, Moscow, Russia. <sup>452</sup>Department of Psychiatry, Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>453</sup>Social Development & Health Promotion Research Center, Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>454</sup>Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>455</sup>Institute of Medicine, University of Colombo, Colombo, Sri Lanka. <sup>456</sup>University of Colombo, Colombo, Sri Lanka. <sup>457</sup>Achutha Menon Centre for Health Science Studies, Sree Chitra Tirunal Institute for Medical Sciences and Technology, Trivandrum, India. <sup>458</sup>Department of Pediatrics & Child Health, Aga Khan University, Karachi, Pakistan. <sup>459</sup>Autism Spectrum Disorders Research Center, Hamadan University of Medical Sciences, Hamadan, Iran. <sup>460</sup>Department of Community Medicine, Banaras Hindu University, Varanasi, India. <sup>461</sup>Manipal Academy of Higher Education, Manipal, India. <sup>462</sup>The George Institute for Global Health, University of New South Wales, New Delhi, India. <sup>463</sup>Environmental Research Center, Duke Kunshan University, Kunshan, China. <sup>464</sup>Nicholas School of the Environment, Duke University, Durham, NC, USA.



<sup>465</sup>Department of Earth Observation Science, University of Twente, Enschede, the Netherlands. <sup>466</sup>Department of Ophthalmology, Heidelberg University, Mannheim, Germany. <sup>467</sup>Beijing Ophthalmology & Visual Science Key Laboratory, Beijing Tongren Hospital, Beijing, China. <sup>468</sup>Department of Community Medicine, Manipal Academy of Higher Education, Mangalore, India. <sup>469</sup>Department of Family Medicine and Public Health, University of Opole, Opole, Poland. <sup>470</sup>School of Health Sciences, Savitribai Phule Pune University, Pune, India. <sup>471</sup>Institute of Family Medicine and Public Health, University of Tartu, Tartu, Estonia. <sup>472</sup>Minimally Invasive Surgery Research Center, Iran University of Medical Sciences, Tehran, Iran. <sup>473</sup>School of Public Health, University College Cork, Cork, UK. <sup>474</sup>Infectious Diseases Research Center, Golestan University of Medical Sciences, Gorgan, Iran. <sup>475</sup>Department of Medical Informatics, Tabriz University of Medical Sciences, Tabriz, Iran. <sup>476</sup>Health Services Management Department, School of Health Qazvin University of Medical Sciences Qazvin, Qazvin, Iran. <sup>477</sup>Community Medicine Department, Rafsanjan University of Medical Sciences, Iran, Rafsanjan, Iran. <sup>478</sup>Department of Forensic Medicine and Toxicology, All India Institute of Medical Sciences, Jodhpur, India. <sup>479</sup>All India Institute of Medical Sciences, New Delhi, India. <sup>480</sup>Department of Epidemiology, Hamadan University of Medical Sciences, Hamadan, Iran. <sup>481</sup>Institute for Epidemiology and Social Medicine, University of Münster, Münster, Germany. <sup>482</sup>Research and Development, Australian Red Cross Blood Service, Sydney, New South Wales, Australia. <sup>483</sup>Hematology-Oncology and Stem Cell Transplantation Research Center, Tehran University of Medical Sciences, Tehran, Iran. <sup>484</sup>Pars Advanced and Minimally Invasive Medical Manners Research Center, Iran University of Medical Sciences, Tehran, Iran. <sup>485</sup>Clinical Pharmacy Unit, Mekelle University, Mekelle, Ethiopia. <sup>486</sup>Department of Anesthesiology & Pain Medicine, University of Washington, Seattle, WA, USA. <sup>487</sup>Department of Public Health, Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>488</sup>Non-Communicable Diseases Research Unit, Medical Research Council South Africa, Cape Town, South Africa. <sup>489</sup>Department of Medicine, University of Cape Town, Cape Town, South Africa. <sup>490</sup>Department of Public Health, Debre Markos University, Debre Markos, Ethiopia. <sup>491</sup>Department of Public Health, Jordan University of Science and Technology, Irbid, Jordan. <sup>492</sup>Social Determinants of Health Research Center, Ahvaz Jundishapur University of Medical Sciences, Ahvaz, Iran. <sup>493</sup>Department of Physiology, Lorestan University of Medical Sciences, Khorramabad, Iran. <sup>494</sup>School of Food and Agricultural Sciences, University of Management and Technology, Lahore, Pakistan. <sup>495</sup>Department of Physiology, Baku State University, Baku, Azerbaijan. <sup>496</sup>School of Health and Rehabilitation Sciences, The University of Queensland, Brisbane, Queensland, Australia. <sup>497</sup>Epidemiology and Biostatistics Department, Health Services Academy, Islamabad, Pakistan. <sup>498</sup>Department of Population Sciences, Jatiya Kabi Kazi Nazrul Islam University, Mymensingh, Bangladesh. <sup>499</sup>Department of Public Health, University of Newcastle, Newcastle, New South Wales, Australia. <sup>500</sup>Department of Hospital Medicine, Miriam Hospital, Brown University, Providence, RI, USA. <sup>501</sup>Department of Internal Medicine, John H. Stroger, Jr. Hospital of Cook County, Chicago, IL, USA. <sup>502</sup>Department of Internal Medicine, Dow University of Health Sciences, Karachi, Pakistan. <sup>503</sup>Faculty of Health and Wellbeing, Sheffield Hallam University, Sheffield, UK. <sup>504</sup>College of Arts and Sciences, Ohio University, Zanesville, OH, USA. <sup>505</sup>Internal Medicine and Gastroenterology Department, National Hepatology and Tropical Research Institute, Cairo, Egypt. <sup>506</sup>Department of Medical Parasitology, Cairo University, Cairo, Egypt. <sup>507</sup>Division of Evidence Synthesis, Datta Meghe Institute of Medical Sciences, Wardha, India. <sup>508</sup>Cancer Research Center, Shahid Beheshti University of Medical Sciences, Tehran, Iran. <sup>509</sup>Academy of Medical Science, Tehran, Iran. <sup>510</sup>Department of Public Health, Mazandaran University of Medical Sciences, Sari, Iran. <sup>511</sup>Department of Biostatistics, Shahid Beheshti University of Medical Sciences, Tehran, Iran. <sup>512</sup>Department of Neurosurgery, Iran University of Medical Sciences, Tehran, Iran. <sup>513</sup>Oxford University Global Surgery Group, University of Oxford, Oxford, UK. <sup>514</sup>Clinical Epidemiology Unit, Lund University, Lund, Sweden. <sup>515</sup>Research and Data Solutions, Synotech Consultant, Nairobi, Kenya. <sup>516</sup>School of Medicine, Xiamen University Malaysia, Sepang, Malaysia. <sup>517</sup>Department of Nutrition, Simmons University, Boston, MA, USA. <sup>518</sup>School of Health Sciences, Kristiania University College, Oslo, Norway. <sup>519</sup>Department of Nursing and Health Promotion, Oslo Metropolitan University, Oslo, Norway. <sup>520</sup>Department of Public Health, Ambo University, Ambo, Ethiopia. <sup>521</sup>Neurophysiology Research Center, Hamadan University of Medical Sciences, Hamadan, Iran. <sup>522</sup>Brain Engineering Research Center, Institute for Research in Fundamental Sciences, Tehran, Iran. <sup>523</sup>Department of Public Health Dentistry, Deemed University, Karad, India. <sup>524</sup>Department of Environmental Health Engineering, Arak University of Medical Sciences, Arak, Iran. <sup>525</sup>Department of Internal and Pulmonary Medicine, Sheri Kashmir Institute of Medical Sciences, Srinagar, India. <sup>526</sup>CIBERSAM, San Juan de Dios Sanitary Park, Sant Boi de Llobregat, Spain. <sup>527</sup>Department of Zoology, University of Oxford, Oxford, UK. <sup>528</sup>Harvard Medical School, Harvard University, Boston, MA, USA. <sup>529</sup>Department of Anthropology, Panjab University, Chandigarh, India. <sup>530</sup>Department of Family and Community Health, University of Health and Allied Sciences, Ho, Ghana. <sup>531</sup>Department of Psychology and Health Promotion, University of KwaZulu-Natal, Durban, South Africa. <sup>532</sup>Department of Psychiatry, University of Nairobi, Nairobi, Kenya. <sup>533</sup>Division of Psychology and Language Sciences, University College London, London, UK. <sup>534</sup>Department of Medicine Brigham and Women's Hospital, Harvard University, Boston, MA, USA. <sup>535</sup>Department of Pathology and Molecular Medicine, McMaster University, Hamilton, Ontario, Canada. <sup>536</sup>Institute of Occupational and Environmental Medicine, University of Birmingham, Birmingham, UK. <sup>537</sup>Health and Nutrition Section, United Nations Children's Fund (UNICEF), Accra, Ghana. <sup>538</sup>Clinical Medicine and Community Health, University of Milan, Milan, Italy. <sup>539</sup>National Institute for Health Research (NIHR), Oxford Biomedical Research Centre, Oxford, UK. <sup>540</sup>Department of Internal Medicine, Post Graduate Institute of Medical Education and Research, Chandigarh, India. <sup>541</sup>Public Health Foundation of India, Gurugram, India. <sup>542</sup>Department of Community and Family Medicine, University of Baghdad, Baghdad, Iraq. <sup>543</sup>School of Medicine, Deakin University, Geelong, Victoria, Australia. <sup>544</sup>Health Promotion and Chronic Disease Prevention Branch, Public Health Agency of Canada, Ottawa, Ontario, Canada. <sup>545</sup>HelpMeSee, New York, NY, USA. <sup>546</sup>International Relations, Mexican Institute of Ophthalmology, Queretaro, Mexico. <sup>547</sup>Department of Otorhinolaryngology (ENT) & Head and Neck Surgery, Father Muller Medical College, Mangalore, India. <sup>548</sup>Department of Information and Internet Technologies, I.M. Sechenov First Moscow State Medical University, Moscow, Russia. <sup>549</sup>Federal Research Institute for Health Organization and Informatics of the Ministry of Health (FRIHOI), Moscow, Russia. <sup>550</sup>School of Nursing, Hong Kong Polytechnic University, Hong Kong, China. <sup>551</sup>School of Pharmacy, Monash University, Bandar Sunway, Malaysia. <sup>552</sup>School of Pharmacy, Taylor's University Lakeside Campus, Subang Jaya, Malaysia. <sup>553</sup>Oxford University Clinical Research Unit, Wellcome Trust Asia Programme, Hanoi, Vietnam. <sup>554</sup>Department of Medicine, University of Malaya, Kuala Lumpur, Malaysia. <sup>555</sup>Department of Medicine and Therapeutics, The Chinese University of Hong Kong, Hong Kong, China. <sup>556</sup>School of Public Health, University of Haifa, Haifa, Israel. <sup>557</sup>Centre for Chronic Disease Control, Beijing, China. <sup>558</sup>Department of Epidemiology, Brown University, Providence, RI, USA. <sup>559</sup>Department of Paediatrics, All India Institute of Medical Sciences, New Delhi, India. <sup>560</sup>Vector Biology, Liverpool School of Tropical Medicine, Liverpool, UK. <sup>561</sup>Department of Nutrition, University of the Philippines Manila, Manila, Philippines. <sup>562</sup>Alliance for Improving Health Outcomes, Inc., Quezon City, Philippines. <sup>563</sup>Institute of Nutrition, Friedrich Schiller University Jena, Jena, Germany. <sup>564</sup>Competence Cluster for Nutrition and Cardiovascular Health (nutriCARD), Jena, Germany. <sup>565</sup>Ariadne Labs, Harvard University, Boston, MA, USA. <sup>566</sup>Development and Communication Studies, University of the Philippines Los Baños, Laguna, Philippines. <sup>567</sup>Pathology Department, Imam Abdulrahman Bin Faisal University, Dammam, Saudi Arabia. <sup>568</sup>Radiology Department, Mansoura University Hospital, Mansoura, Egypt. <sup>569</sup>Ophthalmology Department, Aswan Faculty of Medicine, Aswan, Egypt. <sup>570</sup>Department of Internal Medicine, Grant Medical College & Sir J.J. Group of Hospitals, Mumbai, India. <sup>571</sup>Institute of Medicine, Tribhuvan University, Kathmandu, Nepal. <sup>572</sup>Health Education and Research Department, SDM College of Medical Sciences & Hospital, Dharwad, India. <sup>573</sup>Health University, Rajiv Gandhi University of Health Sciences, Bangalore, India. <sup>574</sup>Environmental Health Research Center, Kurdistan University of Medical Sciences, Sanandaj, Iran. <sup>575</sup>Clinical Research Development Center, Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>576</sup>Department of Maternal and Child Nursing and Public Health, Federal University of Minas Gerais, Belo Horizonte, Brazil. <sup>577</sup>Plastic Surgery Department, Iran University of Medical Sciences, Tehran, Iran. <sup>578</sup>Joint Centre for Bioethics, University of Toronto, Toronto, Ontario, Canada. <sup>579</sup>Ophthalmology Department, Iran University of Medical Sciences, Tehran, Iran. <sup>580</sup>Ophthalmology Department, University of Manitoba, Winnipeg, Manitoba, Canada. <sup>581</sup>School of Science and Health, Western Sydney University, Sydney, New South Wales, Australia. <sup>582</sup>Substance Abuse Prevention Research Center, Kermanshah University of

Medical Sciences, Kermanshah, Iran. <sup>583</sup>Department of Population Studies, University of Zambia, Lusaka, Zambia. <sup>584</sup>Research Department, Grupo de Investigación Fundovida - Fundovida IPS, Cartagena, Colombia. <sup>585</sup>Grupo de Investigación en Economía de la Salud, University of Cartagena, Cartagena, Colombia. <sup>586</sup>Campus Caucaia, Federal Institute of Education, Science and Technology of Ceará, Caucaia, Brazil. <sup>587</sup>Public Health Department, Botho University-Botswana, Gaborone, Botswana. <sup>588</sup>Division of Plastic Surgery, University of Washington, Seattle, WA, USA. <sup>589</sup>Research Department, The George Institute for Global Health, New Delhi, India. <sup>590</sup>School of Medicine, University of New South Wales, Sydney, New South Wales, Australia. <sup>591</sup>ICF International, DHS Program, Rockville, MD, USA. <sup>592</sup>Department of Twin Research and Genetic Epidemiology, King's College London, London, UK. <sup>593</sup>Neurology Department, Janakpuri Super Specialty Hospital Society, New Delhi, India. <sup>594</sup>Neurology Department, Govind Ballabh Institute of Medical Education and Research, New Delhi, India. <sup>595</sup>Pharmacology and Toxicology, Hamadan University of Medical Sciences, Hamadan, Iran. <sup>596</sup>Department of Epidemiology and Biostatistics, University of California San Francisco, San Francisco, CA, USA. <sup>597</sup>Public Health and Mortality, International Institute for Population Sciences, Mumbai, India. <sup>598</sup>Department of Nutrition, University of Oslo, Oslo, Norway. <sup>599</sup>Mekelle University, Mekelle, Ethiopia. <sup>600</sup>Peru Country Office, United Nations Population Fund (UNFPA), Lima, Peru. <sup>601</sup>Forensic Medicine Division, Imam Abdulrahman Bin Faisal University, Dammam, Saudi Arabia. <sup>602</sup>Department of Midwifery, Adigrat University, Adigrat, Ethiopia. <sup>603</sup>Center for Translation Research and Implementation Science, National Institutes of Health, Bethesda, MD, USA. <sup>604</sup>Breast Surgery Unit, Helsinki University Hospital, Helsinki, Finland. <sup>605</sup>University of Helsinki, Helsinki, Finland. <sup>606</sup>Department of Propedeutics of Internal Diseases & Arterial Hypertension, Pomeranian Medical University, Szczecin, Poland. <sup>607</sup>Health Policy and Management, Centre for Regional Policy Research and Cooperation 'Studiorum', Skopje, Macedonia. <sup>608</sup>Pacific Institute for Research & Evaluation, Calverton, MD, USA. <sup>609</sup>Department of Health Research Methods, Evidence and Impact, McMaster University, Hamilton, Ontario, Canada. <sup>610</sup>Global Institute of Public Health (GIPH), Ananthapuri Hospitals and Research Centre, Trivandrum, India. <sup>611</sup>Department of Clinical Biochemistry, Babol University of Medical Sciences, Babol, Iran. <sup>612</sup>Golestan University of Medical Sciences, Gorgan, Iran. <sup>613</sup>Department of Environmental Health, Sabzevar University of Medical Sciences, Sabzevar, Iran. <sup>614</sup>Foodborne and Waterborne Diseases Research Center, Research Institute for Gastroenterology and Liver Diseases, Shahid Beheshti University of Medical Sciences, Tehran, Iran. <sup>615</sup>Kyrgyz State Medical Academy, Bishkek, Kyrgyzstan. <sup>616</sup>Department of Atherosclerosis and Coronary Heart Disease, National Center of Cardiology and Internal Disease, Bishkek, Kyrgyzstan. <sup>617</sup>Research Center for Biochemistry and Nutrition in Metabolic Diseases, Kashan University of Medical Sciences, Kashan, Iran. <sup>618</sup>Department of Rehabilitation and Sports Medicine, Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>619</sup>Deputy of Social Health, Iran University of Medical Sciences, Tehran, Iran. <sup>620</sup>Health Equity Research Center, Tehran University of Medical Sciences, Tehran, Iran. <sup>621</sup>Social Determinants of Health Research Center, Kurdistan University of Medical Sciences, Sanandaj, Iran. <sup>622</sup>Research Center, Salahaddin University, Erbil, Iraq. <sup>623</sup>Internal Medicine Department, King Saud University, Riyadh, Saudi Arabia. <sup>624</sup>Department of Food Technology, Salahaddin University, Erbil, Iraq. <sup>625</sup>Department of Medicine, Karolinska Institutet, Stockholm, Sweden. <sup>626</sup>Department of Information Technology, University of Human Development, Sulaymaniyah, Iraq. <sup>627</sup>Department of Biostatistics, Hamadan University of Medical Sciences, Hamadan, Iran. <sup>628</sup>Department of Epidemiology and Biostatistics, Shahrekord University of Medical Sciences, Shahrekord, Iran. <sup>629</sup>Department of Immunology, Babol University of Medical Sciences, Babol, Iran. <sup>630</sup>Clinical Biochemistry, Tarbiat Modares University, Tehran, Iran. <sup>631</sup>Department of Nursing, Shahrood University of Medical Sciences, Shahrood, Iran. <sup>632</sup>Department of Biomolecular Sciences, University of Mississippi, Oxford, MS, USA. <sup>633</sup>Department of Pharmacy, Mizan-Tepi University, Mizan, Ethiopia. <sup>634</sup>Health Systems and Policy Research Unit, Ahmadu Bello University, Zaria, Nigeria. <sup>635</sup>School of Pharmacy, Haramaya University, Harar, Ethiopia. <sup>636</sup>Iran National Institute of Health Research, Tehran University of Medical Sciences, Tehran, Iran. <sup>637</sup>Community Nutrition, Shahid Beheshti University of Medical Sciences, Tehran, Iran. <sup>638</sup>Health Systems Research Center, National Health Research Institutes, Cuernavaca, Mexico. <sup>639</sup>Department of Public Health Sciences, University of Miami, Miami, FL, USA. <sup>640</sup>Department of Public Health Medicine, University of KwaZulu-Natal, Durban, South Africa. <sup>641</sup>Department of Molecular Medicine, Birjand University of Medical Sciences, Birjand, Iran. <sup>642</sup>Department of Epidemiology and Biostatistics, Kurdistan University of Medical Sciences, Sanandaj, Iran. <sup>643</sup>Department of Epidemiology, Iran University of Medical Sciences, Tehran, Iran. <sup>644</sup>Department of Economics and Management Sciences for Health, Tehran University of Medical Sciences, Tehran, Iran. <sup>645</sup>Department of Mathematical Sciences, University of Bath, Bath, UK. <sup>646</sup>Department of Surgery, University of Washington, Seattle, WA, USA. <sup>647</sup>Department of Clinical Biochemistry, Tarbiat Modares University, Tehran, Iran. <sup>648</sup>Food Science, University of Campinas, Campinas, Brazil. <sup>649</sup>Friedman School of Nutrition Science and Policy, Tufts University, Boston, MA, USA. <sup>650</sup>Federal Institute for Population Research, Wiesbaden, Germany. <sup>651</sup>Center for Population and Health, Wiesbaden, Germany. <sup>652</sup>Indian Institute of Public Health - Hyderabad, Public Health Foundation of India, Hyderabad, India. <sup>653</sup>School of Medical Sciences, Science University of Malaysia, Kubang Kerian, Malaysia. <sup>654</sup>Department of Pediatric Medicine, Nishtar Medical University, Multan, Pakistan. <sup>655</sup>Department of Pediatrics & Pediatric Pulmonology, Institute of Mother & Child Care, Multan, Pakistan. <sup>656</sup>Department of Microbiology and Immunology, Mekelle University, Mekelle, Ethiopia. <sup>657</sup>Department of Urology, Tehran University of Medical Sciences, Tehran, Iran. <sup>658</sup>Department of Medicine, Icahn School of Medicine at Mount Sinai, New York, NY, USA. <sup>659</sup>Research and Analytics, Initiative for Financing Health and Human Development, Chennai, India. <sup>660</sup>Research and Analytics, Bioinsilico Technologies, Chennai, India. <sup>661</sup>Initiative for Non Communicable Diseases, International Centre for Diarrhoeal Disease Research, Dhaka, Bangladesh. <sup>662</sup>Comprehensive Cancer Center, University of Alabama at Birmingham, Birmingham, AL, USA. <sup>663</sup>Department of Epidemiology & Biostatistics, Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>664</sup>Department of Disease, Epidemics, and Pandemics Control, Ministry of Public Health, Yaoundé, Cameroon. <sup>665</sup>Department of Public Health, University of Yaoundé I, Yaoundé, Cameroon. <sup>666</sup>Hospital of the Federal University of Minas Gerais, Federal University of Minas Gerais, Belo Horizonte, Brazil. <sup>667</sup>Department of Pediatrics, Arak University of Medical Sciences, Arak, Iran. <sup>668</sup>Iranian Ministry of Health and Medical Education, Tehran, Iran. <sup>669</sup>General Surgery, Emergency Hospital of Bucharest, Bucharest, Romania. <sup>670</sup>Anatomy and Embryology, Carol Davila University of Medicine and Pharmacy, Bucharest, Romania. <sup>671</sup>Cardiology, Cardio-Aid, Bucharest, Romania. <sup>672</sup>Department of Biological Sciences, University of Embu, Embu, Kenya. <sup>673</sup>Institute for Global Health Innovations, Duy Tan University, Hanoi, Vietnam. <sup>674</sup>Institute of Mental Health Research, University of Ottawa, Ottawa, Ontario, Canada. <sup>675</sup>Department of Clinical Epidemiology, Institute for Clinical Evaluative Sciences, Ottawa, Ontario, Canada. <sup>676</sup>Department of Pharmacology of Tehran University of Medical Sciences, Shahid Beheshti University of Medical Sciences, Tehran, Iran. <sup>677</sup>Heidelberg University Hospital, Heidelberg, Germany. <sup>678</sup>Public Health Department, Universitas Negeri Semarang, Kota Semarang, Indonesia. <sup>679</sup>Graduate Institute of Biomedical Informatics, Taipei Medical University, Taipei City, Taiwan. <sup>680</sup>School of Public Health and Family Medicine, University of Cape Town, Cape Town, South Africa. <sup>681</sup>Department of Neurobiology, Care Sciences and Society (NVS), H1, Division of Family Medicine and Primary Care, Karolinska Institutet, Huddinge, Sweden. <sup>682</sup>Administrative and Economic Sciences, University of Bucharest, Bucharest, Romania. <sup>683</sup>Centre of Cardiovascular Research and Education in Therapeutics, Monash University, Melbourne, Victoria, Australia. <sup>684</sup>Independent Consultant, Accra, Ghana. <sup>685</sup>Department Obstetrics and Gynecology, University of Ibadan, Ibadan, Nigeria. <sup>686</sup>Department of Preventive Medicine, Kyung Hee University, Dongdaemung-gu, South Korea. <sup>687</sup>HAST, Human Sciences Research Council, Durban, South Africa. <sup>688</sup>School of Public Health, University of Namibia, Osakhati, Namibia. <sup>689</sup>Department of Medical Genetics, School of Advanced Technologies in Medicine, Golestan University of Medical Sciences, Gorgan, Iran. <sup>690</sup>Department of Psychiatry and Behavioural Neurosciences, McMaster University, Hamilton, Ontario, Canada. <sup>691</sup>Department of Psychiatry, University of Lagos, Lagos, Nigeria. <sup>692</sup>Centre for Healthy Start Initiative, Lagos, Nigeria. <sup>693</sup>Centre for Healthy Start Initiative, Phonics Hearing Centre, Lagos, Nigeria. <sup>694</sup>Public Health and School of Graduates Studies, Jigjiga University, Jig-Jiga, Ethiopia. <sup>695</sup>Department of Pharmacology and Therapeutics, University of Nigeria Nsukka, Enugu, Nigeria. <sup>696</sup>Department of Psychology, University of Ghana, Accra, Ghana. <sup>697</sup>Graduate School of Public Health, San Diego State University, San Diego, CA, USA. <sup>698</sup>University of Washington, Seattle, WA, USA. <sup>699</sup>University of Port Harcourt, Port Harcourt, Nigeria. <sup>700</sup>School of Medicine, Autonomous University of Madrid, Madrid, Spain. <sup>701</sup>Department of Nephrology and

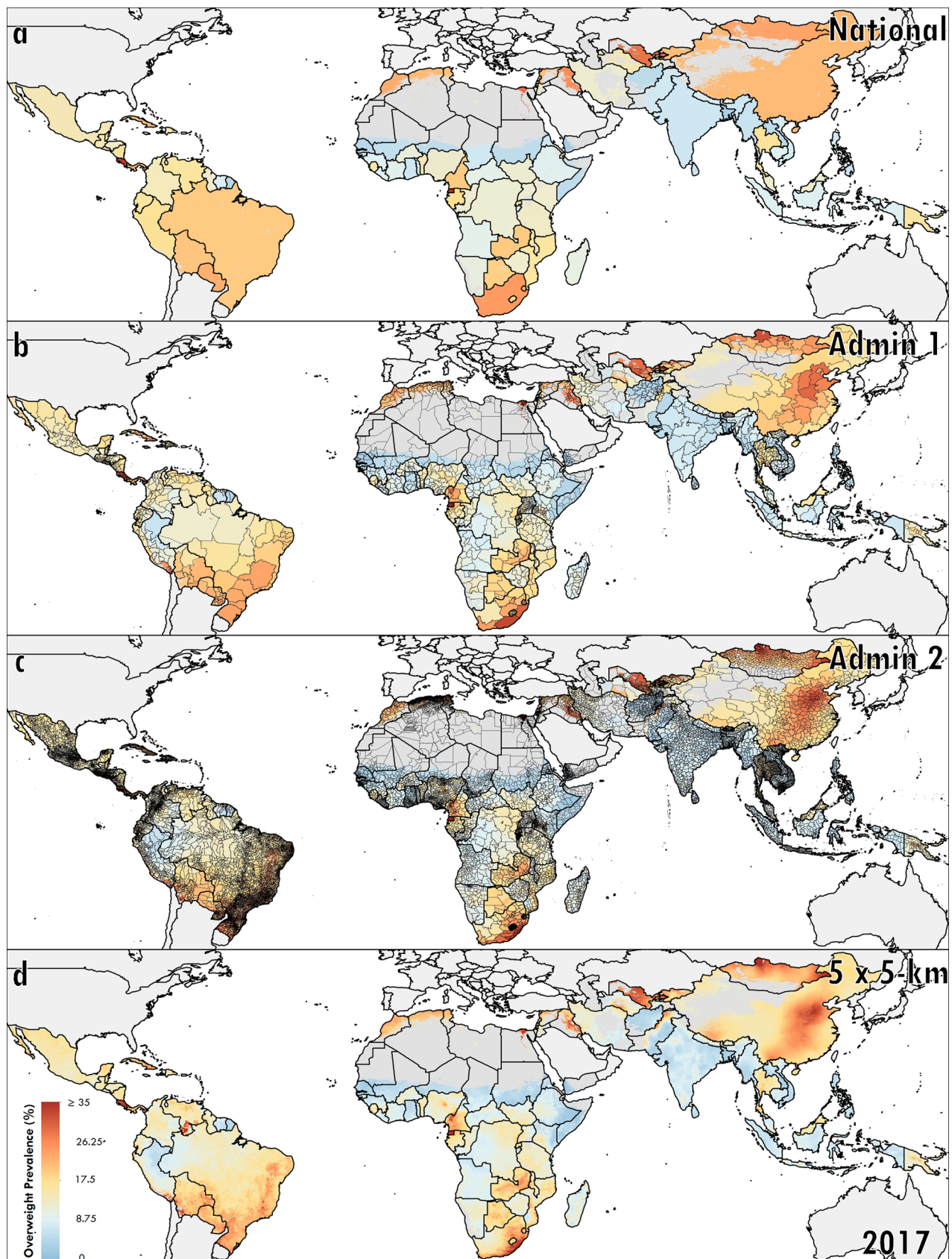
Hypertension, The Institute for Health Research Foundation Jiménez Díaz University Hospital, Madrid, Spain. <sup>702</sup>Department of Environmental Management and Toxicology, University of Benin, Benin City, Nigeria. <sup>703</sup>Institute for Advanced Medical Research and Training, University of Ibadan, Ibadan, Nigeria. <sup>704</sup>Department of Respiratory Medicine, Jagadguru Sri Shivarathreeshwara Academy of Health Education and Research, Mysore, India. <sup>705</sup>Department of Forensic Medicine and Toxicology, Manipal Academy of Higher Education, Mangalore, India. <sup>706</sup>Department of Medical Mycology and Parasitology, Shiraz University of Medical Sciences, Shiraz, Iran. <sup>707</sup>Center for Health Outcomes & Evaluation, Bucharest, Romania. <sup>708</sup>Augenpraxis Jonas, Heidelberg University, Heidelberg, Germany. <sup>709</sup>Internal Medicine, University of Pittsburgh Medical Center, Pittsburgh, PA, USA. <sup>710</sup>Research and Evaluation, Population Council, New Delhi, India. <sup>711</sup>Indian Institute of Health Management Research University, Jaipur, India. <sup>712</sup>Department of Pediatrics, RD Gardi Medical College, Ujjain, India. <sup>713</sup>Public Health Sciences, Karolinska Institutet, Stockholm, Sweden. <sup>714</sup>Research & Publication Cell, Kalinga Institute of Medical Sciences, Bhubaneswar, Bhubaneswar, India. <sup>715</sup>Regional Medical Research Centre, Indian Council of Medical Research, Bhubaneswar, India. <sup>716</sup>Department of Population Studies, International Institute for Population Sciences, Mumbai, India. <sup>717</sup>International Institute of Health Management Research, New Delhi, India. <sup>718</sup>Department of Paediatrics, University of Melbourne, Melbourne, Victoria, Australia. <sup>719</sup>Population Health, Murdoch Childrens Research Institute, Melbourne, Victoria, Australia. <sup>720</sup>Wolaita Sodo University, Sodo, Ethiopia. <sup>721</sup>Department of Physiology, Iran University of Medical Sciences, Tehran, Iran. <sup>722</sup>Center for Research and Innovation, Ateneo De Manila University, Pasig City, Philippines. <sup>723</sup>Istituto di Ricerche Farmacologiche Mario Negri IRCCS, Bergamo, Italy. <sup>724</sup>School of Medicine, University of Virginia, Charlottesville, VA, USA. <sup>725</sup>HIV and Mental Health Department, Integrated Development Foundation Nepal, Kathmandu, Nepal. <sup>726</sup>University Medical Center Groningen, University of Groningen, Groningen, the Netherlands. <sup>727</sup>Faculty of Economics and Business, University of Groningen, Groningen, the Netherlands. <sup>728</sup>Department of Public Health, Maragheh University of Medical Sciences, Maragheh, Iran. <sup>729</sup>Department of Nutrition and Food Sciences, Maragheh University of Medical Sciences, Maragheh, Iran. <sup>730</sup>School of Population and Public Health, University of British Columbia, Vancouver, British Columbia, Canada. <sup>731</sup>Paramedic Department, Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>732</sup>Digestive Diseases Research Institute, Tehran University of Medical Sciences, Tehran, Iran. <sup>733</sup>Fundación Valle del Lili, Cali, Colombia. <sup>734</sup>Infectious Diseases, National Institute of Infectious Diseases, Bucuresti, Romania. <sup>735</sup>Department of Infectious Diseases, Carol Davila University of Medicine and Pharmacy, Bucharest, Romania. <sup>736</sup>Health Sciences Department, Muhammadiyah University of Surakarta, Sukoharjo, Indonesia. <sup>737</sup>Biomedical Engineering Department, Amirkabir University of Technology, Tehran, Iran. <sup>738</sup>Department of Chemistry, Sharif University of Technology, Tehran, Iran. <sup>739</sup>College of Medicine, University of Central Florida, Orlando, FL, USA. <sup>740</sup>College of Graduate Health Sciences, A.T. Still University, Mesa, AZ, USA. <sup>741</sup>Department of Immunology, Mazandaran University of Medical Sciences, Sari, Iran. <sup>742</sup>Molecular and Cell Biology Research Center, Mazandaran University of Medical Sciences, Sari, Iran. <sup>743</sup>Thalassemia and Hemoglobinopathy Research Center, Ahvaz Jundishapur University of Medical Sciences, Ahvaz, Iran. <sup>744</sup>Metabolomics and Genomics Research Center, Tehran University of Medical Sciences, Tehran, Iran. <sup>745</sup>Sina Trauma and Surgery Research Center, Tehran University of Medical Sciences, Tehran, Iran. <sup>746</sup>School of Nursing and Healthcare Professions, Federation University, Heidelberg, Victoria, Australia. <sup>747</sup>National Centre for Farmer Health, Deakin University, Waurn Ponds, Victoria, Australia. <sup>748</sup>Department of Clinical Pediatrics, Sweidi Hospital, Riyadh, Saudi Arabia. <sup>749</sup>Department of Pediatrics, North-West University, Peshawar, Pakistan. <sup>750</sup>Society for Health and Demographic Surveillance, Suri, India. <sup>751</sup>Department of Economics, University of Göttingen, Göttingen, Germany. <sup>752</sup>Birjand University of Medical Sciences, Birjand, Iran. <sup>753</sup>Department of Pharmacology, Shahid Beheshti University of Medical Sciences, Tehran, Iran. <sup>754</sup>University Institute of Public Health, University of Lahore, Lahore, Pakistan. <sup>755</sup>Public Health Department, University of Health Sciences, Lahore, Pakistan. <sup>756</sup>Policy Research Institute, Kathmandu, Nepal. <sup>757</sup>Institute for Poverty Alleviation and International Development, Yonsei University, Wonju, South Korea. <sup>758</sup>Department of Oral Pathology, Srinivas Institute of Dental Sciences, Mangalore, India. <sup>759</sup>Gonçalo Moniz Institute, Oswaldo Cruz Foundation, Salvador, Brazil. <sup>760</sup>Institute of Public Health, Federal University of Bahia, Salvador, Brazil. <sup>761</sup>School of Behavioral Sciences and Mental Health, Tehran Institute of Psychiatry, Tehran, Iran. <sup>762</sup>Kasturba Medical College, Manipal Academy of Higher Education, Mangalore, India. <sup>763</sup>Department of Primary Care and Public Health, Imperial College London, London, UK. <sup>764</sup>Academic Public Health Department, Public Health England, London, UK. <sup>765</sup>WHO Collaborating Centre for Public Health Education and Training, Imperial College London, London, UK. <sup>766</sup>University College London Hospitals, London, UK. <sup>767</sup>School of Health, Medical and Applied Sciences, Central Queensland University, Sydney, New South Wales, Australia. <sup>768</sup>Neurology Department, Sree Chitra Tirunal Institute for Medical Sciences and Technology, Thiruvananthapuram, India. <sup>769</sup>School of Social Sciences and Psychology, Western Sydney University, Penrith, New South Wales, Australia. <sup>770</sup>Translational Health Research Institute, Western Sydney University, Penrith, New South Wales, Australia. <sup>771</sup>Brien Holden Vision Institute, Sydney, New South Wales, Australia. <sup>772</sup>Organization for the Prevention of Blindness, Paris, France. <sup>773</sup>Network of Immunity in Infection, Malignancy and Autoimmunity (NIIMA), Universal Scientific Education and Research Network (USERN), Tehran, Iran. <sup>774</sup>Pediatric Infectious Diseases Research Center, Mazandaran University of Medical Sciences, Sari, Iran. <sup>775</sup>Department of Epidemiology, Birjand University of Medical Sciences, Birjand, Iran. <sup>776</sup>EPIUnit - Public Health Institute University Porto (ISPUP), University of Porto, Porto, Portugal. <sup>777</sup>Surgery Department, University of Minnesota, Minneapolis, MN, USA. <sup>778</sup>Surgery Department, University Teaching Hospital of Kigali, Kigali, Rwanda. <sup>779</sup>School of Psychology, University of Lincoln, Lincoln, UK. <sup>780</sup>Department of Epidemiology and Biostatistics, Imperial College London, London, UK. <sup>781</sup>Department of Clinical Research, Federal University of Uberlândia, Uberlândia, Brazil. <sup>782</sup>Department of Public Health, Wollega University, Nekemte, Ethiopia. <sup>783</sup>Public Health Department, Addis Ababa University, Addis Ababa, Ethiopia. <sup>784</sup>Golestan Research Center of Gastroenterology and Hepatology, Golestan University of Medical Sciences, Gorgan, Iran. <sup>785</sup>Infectious Diseases and Tropical Medicine Research Center, Babol University of Medical Sciences, Babol, Iran. <sup>786</sup>Centro de Investigación Palmira, Agrosavia, Palmira, Colombia. <sup>787</sup>Department of Ocean Science and Engineering, Southern University of Science and Technology, Shenzhen, China. <sup>788</sup>Ain Shams University, Cairo, Egypt. <sup>789</sup>Department of Cardiology, Tehran University of Medical Sciences, Tehran, Iran. <sup>790</sup>National Institute for Research in Environmental Health, Indian Council of Medical Research, Bhopal, India. <sup>791</sup>Cardiovascular Research Institute, Isfahan University of Medical Sciences, Isfahan, Iran. <sup>792</sup>Emergency Department, Shahid Beheshti University of Medical Sciences, Tehran, Iran. <sup>793</sup>Department of Health in Disasters and Emergencies, Shahid Beheshti University of Medical Sciences, Tehran, Iran. <sup>794</sup>Department of Psychiatry, All India Institute of Medical Sciences, New Delhi, India. <sup>795</sup>Halal Research Center of IRI, FDA, Tehran, Iran. <sup>796</sup>Neurogenic Inflammation Research Center, Mashhad University of Medical Sciences, Mashhad, Iran. <sup>797</sup>Nanobiotechnology Center, Soran University, Soran, Iraq. <sup>798</sup>Department of Anatomical Sciences, Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>799</sup>Department of Pathology, Imam Mohammad Ibn Saud Islamic University, Riyadh, Saudi Arabia. <sup>800</sup>Taleghani Hospital, Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>801</sup>Radiology and Nuclear Medicine Department, Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>802</sup>Taleghani Hospital, Kermanshah, Iran. <sup>803</sup>Urology Department, Cairo University, Cairo, Egypt. <sup>804</sup>Public Health and Community Medicine, Cairo University, Giza, Egypt. <sup>805</sup>Drug Applied Research Center, Tabriz University of Medical Sciences, Tabriz, Iran. <sup>806</sup>Department of Entomology, Ain Shams University, Cairo, Egypt. <sup>807</sup>Department of Internal Medicine, University of São Paulo, São Paulo, Brazil. <sup>808</sup>Department of Infectious Diseases and Tropical Medicine, Federal University of Minas Gerais, Belo Horizonte, Brazil. <sup>809</sup>Department of Community Medicine, PSG Institute of Medical Sciences and Research, Coimbatore, India. <sup>810</sup>PSG-FAIMER, South Asia Regional Institute, Coimbatore, India. <sup>811</sup>Health Economics and Financing Research Group, International Centre for Diarrhoeal Disease Research, Bangladesh, Dhaka, Bangladesh. <sup>812</sup>Faculty of Infectious and Tropical Diseases, London School of Hygiene & Tropical Medicine, London, UK. <sup>813</sup>Colorectal Research Center, Iran University of Medical Sciences, Tehran, Iran. <sup>814</sup>Surgery Department, Hamad General Hospital, Hamad Medical Corporation, Doha, Qatar. <sup>815</sup>Faculty of Health & Social Sciences, Bournemouth University, Bournemouth, UK. <sup>816</sup>UGC Centre of Advanced Study in Psychology, Utkal University, Bhubaneswar, India. <sup>817</sup>Udyam-Global Association for Sustainable Development, Bhubaneswar, India. <sup>818</sup>Hypertension in Africa Research Team (HART), North-West University, Potchefstroom, South Africa. <sup>819</sup>Unit for Hypertension and



Cardiovascular Disease, South African Medical Research Council, Cape Town, South Africa. <sup>820</sup>Department of Psychology, University of Alabama at Birmingham, Birmingham, AL, USA. <sup>821</sup>Department of Food Science and Nutrition, Jigjiga University, Jigjiga, Ethiopia. <sup>822</sup>Emergency Department, Manian Medical Centre, Erode, India. <sup>823</sup>Microbiology Service, National Institutes of Health, Bethesda, MD, USA. <sup>824</sup>Department of Health Promotion and Education, Alborz University of Medical Sciences, Karaj, Iran. <sup>825</sup>Health Policy Research Center, Shiraz University of Medical Sciences, Shiraz, Iran. <sup>826</sup>Independent Consultant, Karachi, Pakistan. <sup>827</sup>Department of Neuropsychiatry, Ain Shams University, Cairo, Egypt. <sup>828</sup>School of Medicine, Alborz University of Medical Sciences, Karaj, Iran. <sup>829</sup>Medical Laboratory Sciences, Mazandaran University of Medical Sciences, Sari, Iran. <sup>830</sup>Chronic Diseases (Home Care) Research Center, Hamadan University of Medical Sciences, Hamadan, Iran. <sup>831</sup>Department of Development Studies, International Institute for Population Sciences, Mumbai, India. <sup>832</sup>Department of Basic Sciences, Islamic Azad University, Sari, Iran. <sup>833</sup>Department of Laboratory Sciences, Islamic Azad University, Sari, Iran. <sup>834</sup>University School of Management and Entrepreneurship, Delhi Technological University, New Delhi, India. <sup>835</sup>Department of Health Information Management and Informatics, Iran University of Medical Sciences, Tehran, Iran. <sup>836</sup>Institute for Population Health, King's College London, London, UK. <sup>837</sup>National Institute of Infectious Diseases, Tokyo, Japan. <sup>838</sup>College of Medicine, Yonsei University, Seodaemun-gu, South Korea. <sup>839</sup>Division of Cardiology, Emory University, Atlanta, GA, USA. <sup>840</sup>Finnish Institute of Occupational Health, Helsinki, Finland. <sup>841</sup>Cancer Research Institute, Tehran University of Medical Sciences, Tehran, Iran. <sup>842</sup>Cancer Biology Research Center, Tehran University of Medical Sciences, Tehran, Iran. <sup>843</sup>Institute of Medical Epidemiology, Martin Luther University Halle-Wittenberg, Halle, Germany. <sup>844</sup>Department of Health Education & Promotion, Kermanshah University of Medical Sciences, Kermanshah, Iran. <sup>845</sup>School of Health, University of Technology Sydney, Sydney, New South Wales, Australia. <sup>846</sup>Department of Psychology, Reykjavik University, Reykjavik, Iceland. <sup>847</sup>Department of Health and Behavior Studies, Columbia University, New York, NY, USA. <sup>848</sup>Department of Physical Education, Federal University of Santa Catarina, Florianopolis, Brazil. <sup>849</sup>Department of Law, Economics, Management and Quantitative Methods, University of Sannio, Benevento, Italy. <sup>850</sup>Menzies Institute for Medical Research, University of Tasmania, Hobart, Tasmania, Australia. <sup>851</sup>Global Patient Outcome and Real World Evidence, Eli Lilly and Company, Indianapolis, IN, USA. <sup>852</sup>Department of Humanities and Social Sciences, Indian Institute of Technology, Roorkee, Roorkee, India. <sup>853</sup>Department of Pulmonary Medicine, Asthma Bhawan, Jaipur, India. <sup>854</sup>Department of Medicine, University of Alabama at Birmingham, Birmingham, AL, USA. <sup>855</sup>Medicine Service, US Department of Veterans Affairs, Birmingham, AL, USA. <sup>856</sup>Department of Forensic Medicine, Kathmandu University, Dhulikhel, Nepal. <sup>857</sup>Department of Epidemiology, School of Preventive Oncology, Patna, India. <sup>858</sup>Department of Epidemiology, Healis Sekhsaria Institute for Public Health, Mumbai, India. <sup>859</sup>Department of Midwifery, Haramaya University, Harar, Ethiopia. <sup>860</sup>Department of Physiotherapy and Occupational Therapy, Næstved-Slagelse-Ringsted Hospitals, Slagelse, Denmark. <sup>861</sup>Medical Surgical Nursing Department, Urmia University of Medical Science, Urmia, Iran. <sup>862</sup>Emergency Nursing Department, Semnan University of Medical Sciences, Semnan, Iran. <sup>863</sup>Midwifery Department, Hamadan University of Medical Sciences, Hamadan, Iran. <sup>864</sup>Research Center for Environmental Determinants of Health, Academy of Medical Science, Kermanshah, Iran. <sup>865</sup>Hospital Universitario de la Princesa, Autonomous University of Madrid, Madrid, Spain. <sup>866</sup>Centro de Investigación Biomédica en Red Enfermedades Respiratorias (CIBERES), Madrid, Spain. <sup>867</sup>Department of Research Development, Federal Research Institute for Health Organization and Informatics of the Ministry of Health (FRIHOI), Moscow, Russia. <sup>868</sup>Laboratory of Public Health Indicators Analysis and Health Digitalization, Moscow Institute of Physics and Technology, Moscow, Russia. <sup>869</sup>Hull York Medical School, University of Hull, Hull City, UK. <sup>870</sup>Usher Institute of Population Health Sciences and Informatics, University of Edinburgh, Edinburgh, UK. <sup>871</sup>Department of Parasitology and Mycology, Tabriz University of Medical Sciences, Tabriz, Iran. <sup>872</sup>Division of Community Medicine, International Medical University, Kuala Lumpur, Malaysia. <sup>873</sup>Research Management, Policy, Planning and Coordination, Indian Council of Medical Research, New Delhi, India. <sup>874</sup>Clinical Department, Nutrition and Dietetics Department, Federal Research Institute of Nutrition, Biotechnology and Food Safety, Moscow, Russia. <sup>875</sup>Department of Internal Disease, Pirogov Russian National Research Medical University, Moscow, Russia. <sup>876</sup>Department of Nursing, Muhammadiyah University of Surakarta, Surakarta, Indonesia. <sup>877</sup>Department of Public Health, China Medical University, Taichung City, Taiwan. <sup>878</sup>Department of Community Medicine, Ahmadu Bello University, Zaria, Nigeria. <sup>879</sup>Department of Agriculture and Food Systems, University of Melbourne, Melbourne, Victoria, Australia. <sup>880</sup>Norwegian Institute of Public Health, Bergen, Norway. <sup>881</sup>Department of Community Health, Muhimbili University of Health and Allied Sciences, Dar Es Salaam, Tanzania. <sup>882</sup>Muhimbili University of Health and Allied Sciences, Dar Es Salaam, Tanzania. <sup>883</sup>Department of Criminology, Law and Society, University of California Irvine, Irvine, CA, USA. <sup>884</sup>Department of Medicine, University of Valencia, Valencia, Spain. <sup>885</sup>Carlos III Health Institute, Biomedical Research Networking Center for Mental Health Network (CiberSAM), Madrid, Spain. <sup>886</sup>Cancer Control Center, Osaka International Cancer Institute, Osaka, Japan. <sup>887</sup>Department of Pediatrics, Hawassa University, Hawassa, Ethiopia. <sup>888</sup>International Vaccine Institute, Seoul, South Korea. <sup>889</sup>Research Center for Molecular Medicine, Hamadan University of Medical Sciences, Hamadan, Iran. <sup>890</sup>School of Pharmacy, Mekelle University, Mekelle, Ethiopia. <sup>891</sup>University Institute 'Egas Moniz', Monte da Caparica, Portugal. <sup>892</sup>Research Institute for Medicines, University of Lisbon, Lisbon, Portugal. <sup>893</sup>Department of Public Health, Adigrat University, Adigrat, Ethiopia. <sup>894</sup>Pharmacognosy, Mekelle University, Mekelle, Ethiopia. <sup>895</sup>Department of Pediatrics, King Saud University, Riyadh, Saudi Arabia. <sup>896</sup>College of Medicine, Alfaisal University, Riyadh, Saudi Arabia. <sup>897</sup>Department of Anesthesiology, Perioperative, and Pain Medicine, Stanford University, Stanford, CA, USA. <sup>898</sup>Department of Anesthesiology, King Fahad Medical City, Riyadh, Saudi Arabia. <sup>899</sup>Department of Endocrinology, Christian Medical College and Hospital (CMC), Vellore, India. <sup>900</sup>Biology Department, Moscow State University, Moscow, Russia. <sup>901</sup>HIV/STI Surveillance Research Center, and WHO Collaborating Center for HIV Surveillance, Kerman University of Medical Sciences, Kerman, Iran. <sup>902</sup>Department of Medicine, University of Calgary, Calgary, Alberta, Canada. <sup>903</sup>Department of Pathology and Legal Medicine, University of São Paulo, Ribeirão Preto, Brazil. <sup>904</sup>Clinical Epidemiology and Public Health Research Unit, Burlo Garofolo Institute for Maternal and Child Health, Trieste, Italy. <sup>905</sup>Molecular Medicine and Pathology, University of Auckland, Auckland, New Zealand. <sup>906</sup>Clinical Hematology and Toxicology, Military Medical University, Hanoi, Vietnam. <sup>907</sup>Department of Neurology, All India Institute of Medical Sciences, Delhi, India. <sup>908</sup>Department of Pharmacy, Stamford University Bangladesh, Dhaka, Bangladesh. <sup>909</sup>Gomal Center of Biochemistry and Biotechnology, Gomal University, Dera Ismail Khan, Pakistan. <sup>910</sup>TB Culture Laboratory, Mufti Mehmood Memorial Teaching Hospital Dera Ismail Khan, Dera Ismail Khan, Pakistan. <sup>911</sup>Amity Institute of Biotechnology, Amity University Rajasthan, Jaipur, India. <sup>912</sup>Lifestyle Diseases Research Entity, North-West University, Mmabatho, South Africa. <sup>913</sup>Division of Health Sciences, University of Warwick, Coventry, UK. <sup>914</sup>Department of Epidemiology and Biostatistics, Umeå University, Umeå, Sweden. <sup>915</sup>Argentine Society of Medicine, Buenos Aires, Argentina. <sup>916</sup>Velez Sarsfield Hospital, Buenos Aires, Argentina. <sup>917</sup>Central Research Institute of Cytology and Genetics, Federal Research Institute for Health Organization and Informatics of the Ministry of Health (FRIHOI), Moscow, Russia. <sup>918</sup>Christian Medical College and Hospital (CMC), Vellore, India. <sup>919</sup>UKK Institute, Tampere, Finland. <sup>920</sup>Psychosocial Injuries Research Center, Ilam University of Medical Sciences, Ilam, Iran. <sup>921</sup>National AIDS Control Organisation, Ministry of Health, New Delhi, India. <sup>922</sup>Raffles Neuroscience Centre, Raffles Hospital, Singapore, Singapore. <sup>923</sup>Yong Loo Lin School of Medicine, National University of Singapore, Singapore, Singapore. <sup>924</sup>Community & Family Medicine, All India Institute of Medical Sciences, Bathinda, India. <sup>925</sup>Department of Neurology & Stroke Unit, Sant'Anna Hospital, Como, Italy. <sup>926</sup>Occupational Health Unit, Sant'Orsola Malpighi Hospital, Bologna, Italy. <sup>927</sup>Department of Health Care Administration and Economics, National Research University Higher School of Economics, Moscow, Russia. <sup>928</sup>Department of Global Health and Population, Harvard University, Boston, MA, USA. <sup>929</sup>School of Medicine, University of Belgrade, Belgrade, Serbia. <sup>930</sup>Department of Pediatric Endocrinology, Mother and Child Healthcare Institute of Serbia 'Dr Vukan Cupic', Belgrade, Serbia. <sup>931</sup>Foundation University Medical College, Foundation University, Islamabad, Pakistan. <sup>932</sup>Department of Epidemiology and Biostatistics, Wuhan University, Wuhan, China. <sup>933</sup>Demographic Change and Ageing Research Area, Federal Institute for Population Research, Wiesbaden, Germany. <sup>934</sup>Department of Physical Therapy, Naresuan University, Meung District, Thailand. <sup>935</sup>Department of Psychology and Counselling, University of Melbourne, Melbourne, Victoria, Australia. <sup>936</sup>Department of Medicine, University of Melbourne, St Albans,

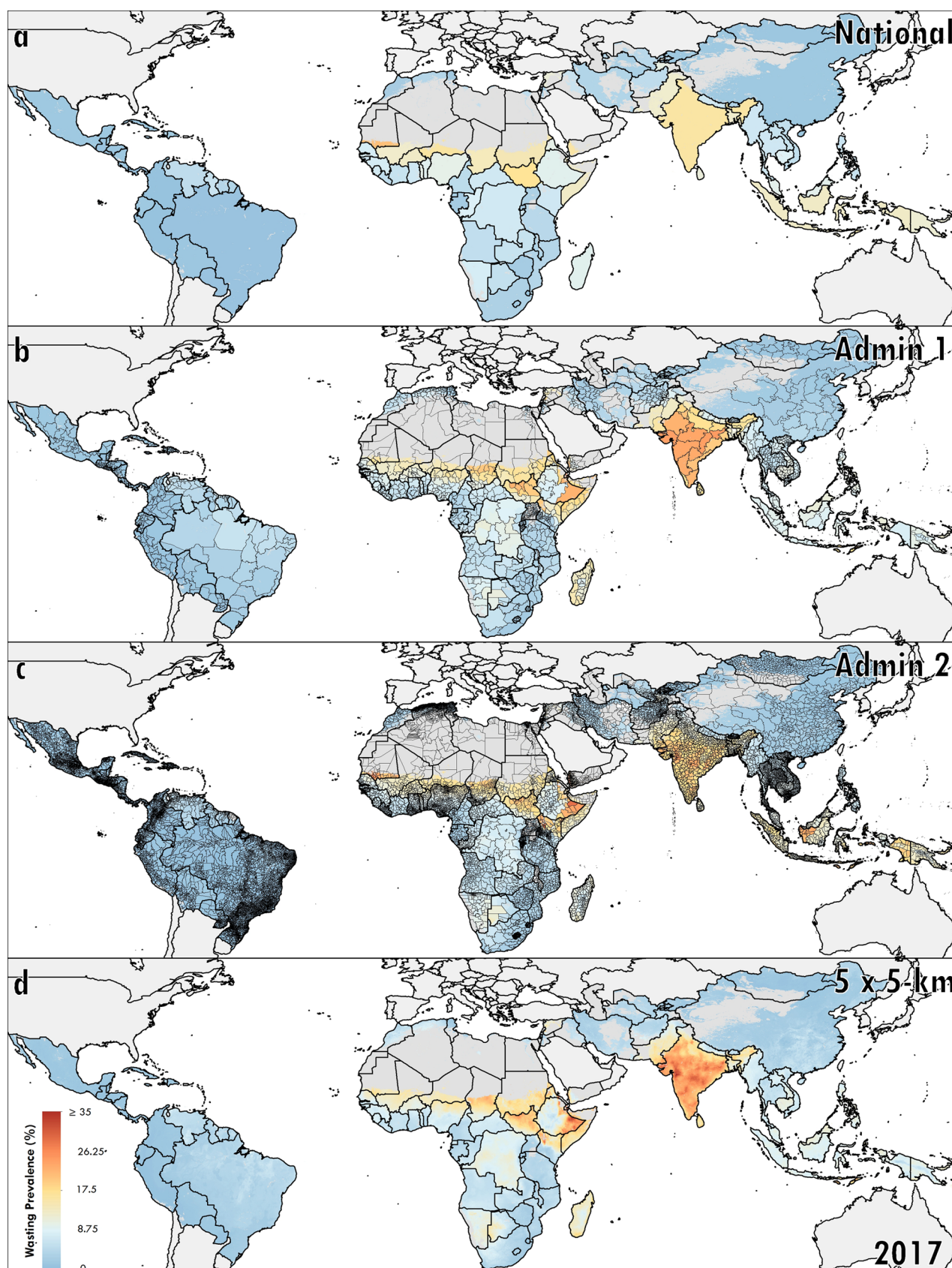


Victoria, Australia. <sup>937</sup>Department of Pharmacology and Toxicology, Mekelle University, Mekelle, Ethiopia. <sup>938</sup>Department of Pharmacology, Addis Ababa University, Addis Ababa, Ethiopia. <sup>939</sup>Department of Nursing, Wollo University, Dessie, Ethiopia. <sup>940</sup>Department of Orthopaedics, Wenzhou Medical University, Wenzhou, China. <sup>941</sup>School of Medicine, Nanjing University, Nanjing, China. <sup>942</sup>Medical Physics Department, Ahvaz Jundishapur University of Medical Sciences, Ahvaz, Iran. <sup>943</sup>Clinical Cancer Research Center, Milad General Hospital, Tehran, Iran. <sup>944</sup>Department of Diabetes and Metabolic Diseases, University of Tokyo, Tokyo, Japan. <sup>945</sup>Department of Preventive Medicine, Northwestern University, Chicago, IL, USA. <sup>946</sup>School of International Development and Global Studies, University of Ottawa, Ottawa, Ontario, Canada. <sup>947</sup>Health Services Management Research Center, Kerman University of Medical Sciences, Kerman, Iran. <sup>948</sup>Department of Health Management, Policy and Economics, Kerman University of Medical Sciences, Kerman, Iran. <sup>949</sup>Wolkite University, Wolkite, Ethiopia. <sup>950</sup>Centre for Suicide Research and Prevention, University of Hong Kong, Hong Kong, China. <sup>951</sup>Department of Social Work and Social Administration, University of Hong Kong, Hong Kong, China. <sup>952</sup>Department of Psychopharmacology, National Center of Neurology and Psychiatry, Tokyo, Japan. <sup>953</sup>Department of Preventive Medicine, Korea University, Seoul, South Korea. <sup>954</sup>Department of Sociology, Yonsei University, Seoul, South Korea. <sup>955</sup>Department of Health Policy & Management, Jackson State University, Jackson, MS, USA. <sup>956</sup>School of Medicine, Tsinghua University, Beijing, China. <sup>957</sup>Department of Environmental Health, Mazandaran University of Medical Sciences, Sari, Iran. <sup>958</sup>Environmental Health, Academy of Medical Science, Sari, Iran. <sup>959</sup>Global Health Institute, Wuhan University, Wuhan, China. <sup>960</sup>Social Determinants of Health Research Center, Ardabil University of Medical Science, Ardabil, Iran. <sup>961</sup>Department of Medicine, Monash University, Melbourne, Victoria, Australia. <sup>962</sup>Student Research Committee, Babol University of Medical Sciences, Babol, Iran. <sup>963</sup>Department of Community Medicine, Ardabil University of Medical Science, Ardabil, Iran. <sup>964</sup>Psychiatry and Psychology Research Center, Tehran University of Medical Sciences, Tehran, Iran. <sup>965</sup>Maternal and Child Wellbeing Unit, African Population Health Research Centre, Nairobi, Kenya. <sup>966</sup>Public Health Department, Dilla University, Dilla, Ethiopia. <sup>967</sup>School of Public Health, Wuhan University of Science and Technology, Wuhan, China. <sup>968</sup>Hubei Province Key Laboratory of Occupational Hazard Identification and Control, Wuhan University of Science and Technology, Wuhan, China. <sup>969</sup>Department of Preventive Medicine, Wuhan University, Wuhan, China. <sup>970</sup>School of Biology and Pharmaceutical Engineering, Wuhan Polytechnic University, Wuhan, China. ✉e-mail: [sihay@uw.edu](mailto:sihay@uw.edu)



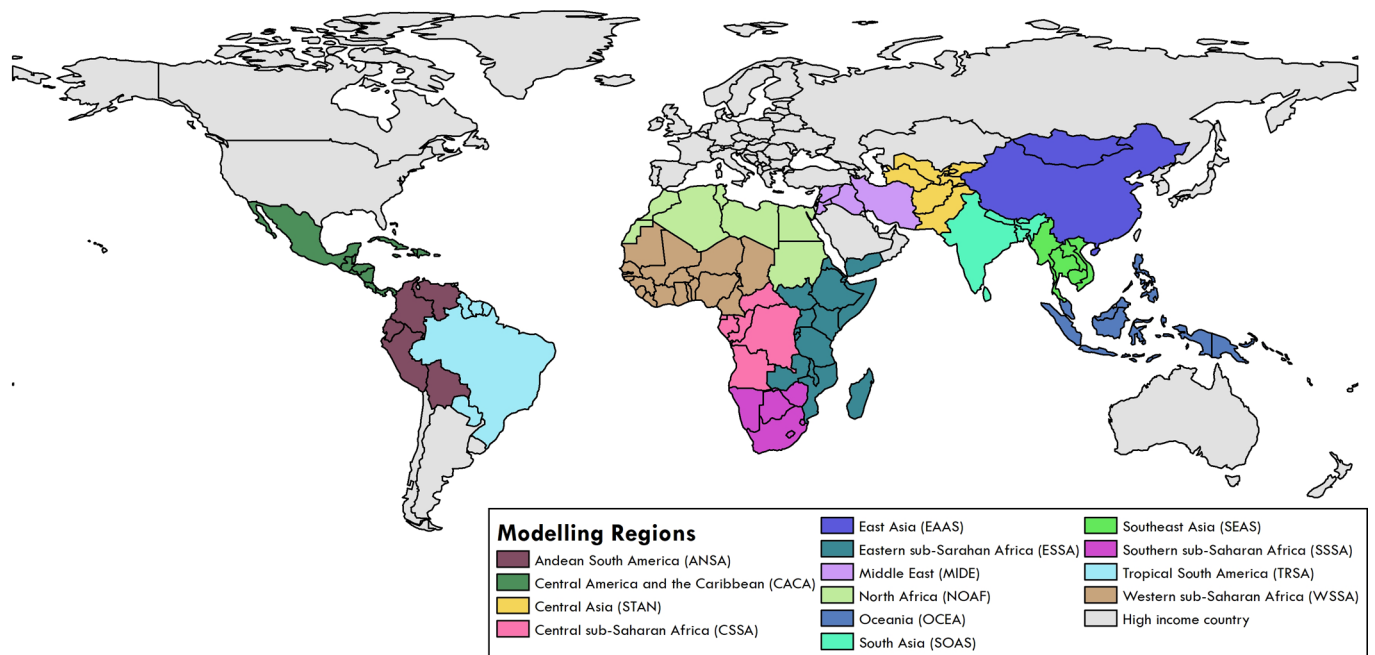
**Extended Data Fig. 1 | Prevalence of under-5 childhood overweight in LMICs in 2017 at administrative levels 0, 1, 2, and at 5 × 5-km resolution.**

Prevalence of overweight among children under 5 at administrative level 0 (national-level estimates) (a), first administrative unit (b), second administrative unit (c), and at the 5 × 5-km resolution (d). Maps reflect administrative boundaries, land cover, lakes, and population; grey-coloured grid cells were classified as "barren or sparsely vegetated" and had fewer than ten people per 1 × 1-km grid cell<sup>39–45</sup>, or were not included in this analysis. Maps were generated using ArcGIS Desktop 10.6.



**Extended Data Fig. 2 | Prevalence of under-5 child wasting in LMICs at administrative levels 0, 1, 2, and at 5 × 5-km resolution in 2017.** Prevalence of wasting among children under 5 at administrative level 0 (national-level estimates) (a), first administrative unit (b), second administrative unit (c), and at the 5 × 5-km resolution (d). Maps reflect administrative boundaries, land cover, lakes, and population; grey-coloured grid cells were classified as “barren or sparsely vegetated” and had fewer than ten people per 1 × 1-km grid cell<sup>39–45</sup>, or were not included in this analysis. Maps were generated using ArcGIS Desktop 10.6.





**Extended Data Fig. 3 | Modelling regions.** Modelling regions<sup>46</sup> were based on geographic and socio-demographic index (SDI) regions from the Global Burden of Disease<sup>47</sup>, defined as: Andean South America, Central America and the Caribbean, Central sub-Saharan Africa (SSA), East Asia, Eastern SSA, Middle East, North Africa, Oceania, Southeast Asia, South Asia, South SSA, Central Asia, Tropical South America, and Western SSA. Regions in grey (Stage 3) were not included in our models due to high-middle and high SDI. Map was generated using ArcGIS Desktop 10.6.

46. Murray, C. J. et al. GBD 2010: design, definitions and metrics. *Lancet* **380**, 2063–2066 (2012).

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| <input type="checkbox"/>            | <input checked="" type="checkbox"/> A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons  |
| <input type="checkbox"/>            | <input checked="" type="checkbox"/> A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals) |
| <input checked="" type="checkbox"/> | <input type="checkbox"/> For null hypothesis testing, the test statistic (e.g. $F$ , $t$ , $r$ ) with confidence intervals, effect sizes, degrees of freedom and $P$ value noted<br><i>Give <math>P</math> values as exact values whenever suitable.</i>                                       |
| <input type="checkbox"/>            | <input checked="" type="checkbox"/> For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings   |
| <input type="checkbox"/>            | <input checked="" type="checkbox"/> For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes   |
| <input checked="" type="checkbox"/> | <input type="checkbox"/> Estimates of effect sizes (e.g. Cohen's $d$ , Pearson's $r$ ), indicating how they were calculated  |

Our web collection on [statistics for biologists](#) contains articles on many of the points above.

### Software and code

Policy information about [availability of computer code](#)

Data collection No primary data collection was carried out for this analysis

Data analysis This analysis was carried out using R version 3.5.0. The main geostatistical models were fit using R-INLA version 18.07.12. **Additional adjustments were performed using the mgcv package in R (v. 3.5.0).** All code used for these analyses is publicly available online at <http://ghdx.healthdata.org/>. Maps were generated using ArcGIS Desktop 10.6.

For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors/reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Research [guidelines for submitting code & software](#) for further information.

### Data

Policy information about [availability of data](#)

All manuscripts must include a [data availability statement](#). This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A list of figures that have associated raw data
- A description of any restrictions on data availability

The findings of this study are supported by data available in public online repositories, data that are publicly available upon request from the data provider, and data that are not publicly available due to restrictions by the data provider and which were used under license for the current study. A detailed table of data sources and availability can be found in Supplementary Table 2, and online at [ghdx.healthdata.org](http://ghdx.healthdata.org).

## Field-specific reporting

Please select the one below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.

☒ Life sciences ☐ Behavioural & social sciences ☐ Ecological, evolutionary & environmental sciences

For a reference copy of the document with all sections, see [nature.com/documents/nr-reporting-summary-flat.pdf](https://www.nature.com/documents/nr-reporting-summary-flat.pdf)

## Life sciences study design

All studies must disclose on these points even when the disclosure is negative.

Sample size	Sample size was calculated as the number of unique data source-location pairs with observations of overweight and wasting prevalence. This sample size is reported in the main text under Global and location variation in malnutrition trends, "...using data from 420 household surveys representing more than 3 million children, we map the relative burdens of overweight and wasting among under-5 children in 105 LMICs from 2000 to 2017."
Data exclusions	Reasons for data exclusion were pre-established and are described in supplementary table 5. For a survey to be considered for this analysis, we required information on height, weight, age and sex. Select data sources were excluded from the analysis due to: missing survey weights, missing sex and age variable, incomplete sampling (e.g., only a specific age range), or untrustworthy data (as determined by the survey administrator or by inspection).
Replication	This is an observational study using many years of survey and surveillance data and could be replicated.
Randomization	This analysis is an observational mapping study and there were no experimental groups.
Blinding	Blinding was not relevant to this study, as it was an observational study using survey and surveillance data.

## Reporting for specific materials, systems and methods

We require information from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, system or method listed is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.

### Materials & experimental systems

### Methods

n/a	Involved in the study	n/a	Involved in the study
<input checked="" type="checkbox"/>	<input type="checkbox"/> Antibodies	<input checked="" type="checkbox"/>	<input type="checkbox"/> ChIP-seq
<input checked="" type="checkbox"/>	<input type="checkbox"/> Eukaryotic cell lines	<input checked="" type="checkbox"/>	<input type="checkbox"/> Flow cytometry
<input checked="" type="checkbox"/>	<input type="checkbox"/> Palaeontology	<input checked="" type="checkbox"/>	<input type="checkbox"/> MRI-based neuroimaging
<input checked="" type="checkbox"/>	<input type="checkbox"/> Animals and other organisms		
<input checked="" type="checkbox"/>	<input type="checkbox"/> Human research participants		
<input checked="" type="checkbox"/>	<input type="checkbox"/> Clinical data		